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RESULT 946
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                    ABT11934
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention provides a composition comprising one or more antisense oligonucleotides directed against vascular endothelial growth factor (VEGF) where the antisense oligonucleotides inhibits proliferation of cells exhibiting autoorine VEGF activity at an IC 50 concentration of between 0.5-2.5 micro Ma. The antisense oligonucleotides may be directed against VEGF for inhibiting cancer cell proliferation and angiogenesis. Preferably the oligonucleotide AMI23032 (a modified version of AAMI23984) is used and may be utilized to treat Kaposi's sarcoma, ovarian cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    prostrate cancer, pancreatic cancer or melanoma. Sequences AAH23012-023 represent gene-specific primers used in RT-PCR amplification of VEGF
                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel antisense oligonucleotides useful for inhibiting vascular endothelial growth factor expression, angiogenesis and for treating cancer, e.g., Kaposi's sarcoma, ovarian cancer and prostrate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                        Vascular endothelial growth factor; VEGF; antisense; angiogenesis; cell proliferation; Kaposi's sarcoma; cancer; melanoma; cytostatic; antisense therapy; RT-PCR; primer; VEGFR-2; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03; tive . 0; Mismatches 3; Indels
                                                                0.4%; Score 16.2; DB 1; Length 23;
85.7%; Pred. No. 1.4e+03;
ive 0; Mismatches 3; Indels
                                  Sequence 23 BP; 6 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 4 A; 7 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                     VEGFR-2 gene specific reverse primer
                                                                                                                                         3185 TCCAGCCTGCCCCGGAGCTGG 3205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 12; Page 56; 105pp; English
                                                                                                                                                                          rccarcrrccccccaccrcc 2
the zeta-COP nucleotide sequence
                                                                                                                                                                                                                                                                ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JAN-2001; 2001WO-US000019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-JAN-2000; 2000US-00487023
                                                                                                                                                                                                                                                                AAH23021 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                    Local Similarity 85.7
1es 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-451898/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gill PS, Masood R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GILL/) GILL P.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200152904-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                      17-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                        22
                                                                     Query Match
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                                                                                                                                                                                                                             RESULT 945
                                                                                      Best Loc
Matches
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The invention comprises a transgenic ungulate containing a xenogenous finamuooplobulin gene which undergoes rearrangement and expresses more than one xenogenous immunoglobulin molecule. The transgenic ungulates of the invention are useful for producing antibodies that are useful for producing antibodies that are useful for propplylaxis/traemment of infectious diseases, modulation of the immune system, removal of cancer cells and modulation of specific human molecules. The present DNA sequence represents a PCR primer that was used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New transgenic ungulates comprising nucleic acids encoding a xenogenous immunoglobulin (Ig) that undergoes rearrangement and expresses xenogenous Ig molecules, useful for producing antibodies for prophylaxis of infectious diseases.
                                                                                                                                                                                                            Human, PCR, primer; ss, transgenic ungulate, xenogenous immunoglobulin; infectious disease; cancer cell removal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tomizuka K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kuroiwa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunoglobulin heavy chain PCR primer #14.
                                                                                                                                                          Human immunoglobulin heavy chain PCR primer #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ferguson SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 59; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           854 AGGAGGAGCTGGTGGAGGCTG 874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGGTGCAGCTGGTGGAGTCTG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in an example of the invention
ABT11934 standard; DNA; 23 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-NOV-2000; 2000US-00714185.
20-DEC-2000; 2000US-0256458P.
09-AUG-2001; 2001US-0311625P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-NOV-2001; 2001WO-US043128
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                                                                                                       19-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AURO-) AUROX LLC.
(KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Goldsby RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-698746/75.
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                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                        12-SEP-2002.
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Ishida I;
                                                     ABT11934;
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Matches
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Gaps

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1950 GATCATGCGGGAGTGCTGGCA 1970

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Best Local Similarity 85.74 Matches 18; Conservative

21 GACCATGCTGGACTGCTGGCA 1

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The invention comprises a transgenic ungulate containing a xenogenous immunoglobulin gene which undergoes rearrangement and expresses more than one xenogenous immunoglobulin molecule. The transgenic ungulates of the invention are useful for producing antibodies that are useful for prophylaxis/treatment of infectious diseases, modulation of the immune system, removal of cancer cells and modulation of specific human molecules. The present DNA sequence represents a PCR primer that was used in an example of the invention
                                                                                                                   New transgenic ungulates comprising nucleic acids encoding a xenogenous immunoglobulin (Ig) that undergoes rearrangement and expresses xenogenous Ig molecules, useful for producing antibodies for prophylaxis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant DNA construct; secretory leader sequence; His tag; infection; therapy; RGDP; secreted replicable genetic display package; vaccination; restriction endonuclease; cleavage site; phage display library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a recombinant DNA construct coding for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel recombinant DNA construct comprising DNA sequence for secreted replicable genetic display package out of frame with DNA sequences for secretory leader sequence, tag, and two restriction enzyme recognition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer HuVH3aBACK used to generate human scFv antibodies.
                                 Tomizuka K;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.2; DB 1; Length 23; Pred. No. 1.4e+03; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                  Kuroiwa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibacterial; PCR primer; human; ss.
                                  Ferguson SE,
                                                                                                                                                                                                               Example 2; Page 60; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 31; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 854 AGGAGGACCTGGTGGAGGCTG 874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 Agerecaecreerecaerere 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rigg G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-AUG-2000; 2000GB-00019049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD30849 standard; DNA; 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 85.7
Matches 18; Conservative
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(KIRI ) KIRIN BEER KK
                                    Goldsby RA,
                                                                                                                                                                              infectious diseases.
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                                                                                        WPI; 2002-698746/75
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                                  Robl JM,
Ishida I;
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                                 Robl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention comprises a transgenic ungulate containing a xenogenous immunoglobulin gene which undergoes rearrangement and expresses more than one xenogenous immunoglobulin molecule. The transgenic ungulates of the invention are useful for producing antibodies that are useful for prophylaxis/treatment of infectious diseases, modulation of the immune system, removal of cancer cells and modulation of specific human molecules. The present DNA sequence represents a PCR primer that was used in an example of the invention
                                                                                                                                                                                                                                                                                                                                                       New transgenic ungulates comprising nucleic acids encoding a xenogenous immunoglobulin (Ig) that undergoes rearrangement and expresses xenogenous Ig molecules, useful for producing antibodies for prophylaxis of infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; PCR; primer; ss; transgenic ungulate; xenogenous immunoglobulin; infectious disease; cancer cell removal.
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                                                                                                                                                                                                                                                                    Tomizuka K;
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85.7%; Pred. No. 1.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                    Kuroiwa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunoglobulin heavy chain PCR primer #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                      Ferguson SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 63; 132pp; English.
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20-DEC-2000; 2000US-0256458P.
09-AUG-2001; 2001US-0311625P.
                                                                                                                                         17-NOV-2000; 2000US-00714185.
20-DEC-2000; 2000US-0256458P.
09-AUG-2001; 2001US-0311625P.
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                                                                                                            16-NOV-2001; 2001WO-US043128
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les 18; Conservative
                                                                                                                                                                                                                                                                      Goldsby RA,
                                                                                                                                                                                                                 (AURO-) AUROX LLC.
(KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                      WPI; 2002-698746/75.
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                                      WO200270648-A2
       Homo sapiens.
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                                                                       12-SEP-2002.
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                                                                                                                                                                                                                                                                    Robl JM,
Ishida I;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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g 8

Secretory leader sequence, tag, at least two restriction endonuclease cleavage sites, and a component of a secreted replicable genetic display package (RGDP) which, when expressed in frame in a recombinant host craganism, causes the RGDP to display at the surface of the package, the tag and any polypeptide encoded 3' of the tag. The method of the corganism, causes the RGDP to display at the surface of the package, the invention is also useful for producing a member of SBP (specific binding pairs) where the recombinant host construct is expressed in recombinant to host organism or recombinant host cells. The recombinant host organism or constructs prepared with a helper phage and the method involves isolating the RGDPs displaying the tag. The method is useful for manufacturing a phage displaying the tag. The method is useful for manufacturing a phage displayl library where several recombinant DNA constructs prepared using several DNA sequences for components of specific binding pairs, are employed. Members (antibodies or their antigen binding pairs, are employed. Members (antibodies or their antigen binding fragments, or ScPv fragments) of SBP identified have therapeutic or diagnostic applications. The coding sequence may be determined and synthesised in bulk for use as diagnostic or therapeutic agent for treating infection. The SBP members are also useful for isolating and purifying their complementary binding partners which are useful of sequences are used for preparing a library of databases of sequences of sequences are used for preparing a library of databases of sequences are useful for fragment. This library can be examined to determine the identity of the antigen binding fragments are useful for fragment. This library can be examined to determine the identity of the antigen bound by the antibody or its fragment. This great is a possible and antigen beneficed to determine the identity of the antigen beneficed in the generation of human scfv antibodies 88888888888888888888888888888888888

Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;

0; Gaps Query Match 0.4%; Score 16.2; DB 1; Length 23; Best Local Similarity 85.7%; Pred. No. 1.4e+03; Matches 18; Conservative 0; Mismatches 3; Indels 3; Indels

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ઠે 원 RESULT 950

ACA94758 standard; DNA; 23

BP

21-JUL-2003 (first entry) ACA94758;

Human single chain variable fragment PCR primer #2.

Single chain monoclonal antibody; cytostatic; virucide; anti-HIV; hepatchropic; antiinflammatory; gene therapy; DNA construct library; transcription; transcriptional activator; antigen binding domain; immunoglobulin; endogenous transcriptional associated regulatory protein; affinity purification; drug screening; physiological disorder; cancer; viral infection; hepatitis; respiratory syncytial virus; RSV; HIV; wiral infection; intus; respiratory syncytial virus; RSV; HIV; measles; dengue virus; Ebola virus; PCR; primer; ss.

Homo sapiens.

US2003017149-A1.

23-JAN-2003

28-AUG-2001; 2001US-00939769

10-OCT-1996;

(HOEF/) HOEFFLER J P. (RUSS/) RUSSELL M. Hoeffler JP, Russell M;

WPI; 2003-428947/40.

Screening a DNA construct library for single chain monoclonal antibody fusion reagents for regulating transcription in vivo, by using constructs expressing antigen binding domains that target transcriptional activators.

Claim 23; Page 15; 52pp; English

The invention describes a method of screening a DNA construct library for a single chain monoclonal antibody fusion reagent capable of binding a transcriptional associated biomolecule in vivo comprising employing the activity of a transcriptional activator by using constructs that express antigen binding domains of immunoglobulins to target endogenous cranscriptional associated regulatory proteins The method is useful for screening a DNA construct library for single chain monoclonal antibody fusion reagents capable of binding a transcriptional associated content in vivo, or capable of regulating transcription in vivo. The identified single chain monoclonal antibody fusion reagents are useful in affinity purification, drug screening, gene therapy (particularly for contenting the transcription of a gene in vivo), or in diagnosing a physiological disorder manifested by abnormal levels of transcription associated biomolecules. The single chain monoclonal antibody fusion cagents may be used for treating or diagnosing cancer or viral infections (e.g. hepatitis A or B, respiratory syncytial virus (RSV), HV, Junin virus, herpes simplex virus, rubella, Varicella-Zoster virus, measles, dengue virus or Ebola virus infections). This sequence represents a primer used to isolate human single chain variable fragment library

Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;

Gaps Indels 0; Query Match 0.4%; Score 16.2; DB 1; Length 23; Best Local Similarity 85.7%; Pred. No. 1.4e+03; Matches 18; Conservative 0; Mismatches 3; Indels

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854 AGGAGGAGCTGGTGGAGGCTG 874 2 AGGTGCAGCTGGTGAGTCTG 22 셤 ઠે

RESULT 951 ACA94762

ACA94762 standard; DNA; 23 BP

ACA94762;

21-JUL-2003 (first entry)

Human single chain variable fragment PCR primer #6.

Single chain monoclonal antibody, cytostatic; virucide; anti-HIV; hepatchropic; antiinflammatory, gene therapy, DNA construct library; transcription; transcriptional activator; antigen binding domain; immunoglobulin; endogenous transcriptional associated requlatory protein; affinity purification; drug screening; physiological disorder; cancer; viral infection; hepatitis; respiratory syncytial virus; RSV; HIV; measles; dengue virus; Ebola virus; PCR; primer; ss.

Homo sapiens.

US2003017149-A1.

23-JAN-2003.

28-AUG-2001; 2001US-00939769.

96US-00728890. 10-OCT-1996;

(HOEF/) HOEFFLER J P. (RUSS/) RUSSELL M.

Russell

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RIPE3b1 peptide-based 3' PCR primer.
                                              Disclosure; Page 15; 52pp; English
                                                                                                                                                                                                        854 AGGAGGAGCTGGTGGAGGCTG 874
                                                                                                                                                                                                                  Acerecaecreerecaerere 22
                                                                                                                                                                                                                                                                                                                                                             (JOSL-) JOSLIN DIABETES CENT INC.
                                                                                                                                                                                                                                                                                                                                                   31-AUG-2001; 2001US-0316453P.
                                                                                                                                                                                                                                                                                                                                        30-AUG-2002; 2002WO-US027600
                                                                                                                                                                                                                                            ACC48959 standard; DNA; 23
                                                                                                                                                                                             18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-300877/29
          WPI; 2003-428947/40
                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                    WO2003020894-A2.
                                                                                                                                                                                                                                                                                                          Cricetulus sp.
                                                                                                                                                                                                                                                                 11-AUG-2003
                                                                                                                                                                                                                                                                                                                              13-MAR-2003.
                                                                                                                                                                                                                                                                                               primer; ss.
                                     activators
                                                                                                                                                                                                                                                                                                                                                                        Sharma A;
                                                                                                                                                                                                                                                      ACC48959;
                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                  952
                                                                                                                                                                                             Matches
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BP.

(first entry)

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The present sequence is a 3' primer based on a RIPEIb1 PCR product obtained from hamster insulinoma HIT T-15 cells. The primer was used in the cloning of human mmdafa CDNA (see Acc19953). mmdafA cell ABR42043) is a novel insulin gene transcription factor. A claimed method of treating an insulin-related disorder comprises increasing the level, expression or activity of a Maf protein, especially mmdafA, in a subject. The insulin-related disorder is diabetes, hypertension, retinopathy, persistent hyperinsulinaemic hypoglycaemia of infancy (PHHI), insulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel method for the identification of candidate sequences of at least the complimentarity-determining region 3 (CDR3) of antibodies specific against at least 1 antigen produced by a microorganism during an infection or against a vaccine. The method comprises initially the sequencing at least the complementarity
Novel isolated mammalian insulin related transcription factor polypeptide, mMafA, useful for treating an insulin-related disorder in a subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying candidate sequences of antibodies specific against at least antigen produced by a microorganism during an infection or against a vaccine, useful;for treating e.g. HIV, bacterial, viral, or parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human complimentarity-determining region 3; CDR3; microorganism;
infection; vaccine; heavy chain variable region; VH; parasitic;
light chain variable region; VL; B cell; antibacterial; virucidal;
fungicidal; anti-HIV; antiparasitic; antibody; HIV; bacterial; viral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer HuVHJaBACK related to human VH and VL antibody fragments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 16.2; DB 1; Length 23; , 85.7%; Pred. No. 1.4e+03; .ive 0; Mismatches 3; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23 BP; 3 A; 12 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ä
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                                                                                                                               Example 6; Page 137; 170pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2103 CACCCCCAGCTCCAGCTCCTC 2123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 38; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Burnie JP, Matthews RC, Rigg GP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 caccrccagcircagciacic 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-DEC-2001; 2001GB-00030267.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABT43625 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NEUT-) NEUTEC PHARMA PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 85.7
Matches 18, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      yeast; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-523552/49.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003052416-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABT43625;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
                                                                                                                                                                                                                                                                                     The invention describes a method of screening a DNA construct library for a single chain monoclonal antibody fusion reagent capable of binding a crinityity of a transcriptional activator by using constructs that express antigen binding domains of immunoglobulins to target endogenous cranscriptional associated regulatory proteins. The method is useful for screening a DNA construct library for single chain monoclonal antibody biomolecule in vivo, or capable of regulating transcriptional antibody biomolecule in vivo, or capable of regulating transcription in vivo. The identified single chain monoclonal antibody fusion reagents associated capable of regulating transcription in vivo. The identified single chain monoclonal antibody fusion reagents are useful in segulating the transcription of a gene therapy (particularly for regulating the transcription of a gene therapy (particularly for regulating the transcription of a gene in vivo), or in diagnosing a physiological disorder manifested by abnormal levels of transcription associated biomolecules. The single chain monoclonal antibody fusion associated biomolecules are simple chain monoclonal antibody fusion infections (e.g. hepatitis A or B, respiratory syncytial virus (RSV), HV, Junin virus, herpes simplex virus, rubella, Varicella-Zoster virus, represents a primer used to isolate human single chain antibody variable fragments for used in the creation of a human single chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                      Screening a DNA construct library for single chain monoclonal antibody fusion reagents for regulating transcription in vivo, by using constructs expressing antigen binding domains that target transcriptional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mMafA; mammalian MafA; transcription factor; RIPE3bl; insulin; diabetes; antidiabetic; hypotensive; ophthalmological; hypoglycaemic; hamster; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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determining region 3 (CDR3) of the heavy chain variable (VH) and/or light chain variable (VL) coding regions of B cells isolated from at least one patient who has been infected by the microorganism or administered the vaccine. Subsequently, the method comprise correlating the sequences of the CDR3 regions of the VH and VL coding regions of the B cells to identify a set of candidate sequences for at least a CDR3 region of antibodies specific against the antigen produced by the microorganism or antibodies specific and the method of the invention may provide agents with antibodies through use as a vaccine. The antibodies may therefore be useful for manufacturing a medicament for treating an infection caused by a microorganism (for example HIV, bacterial, viral, parasitic or yeast infection), for determining the efficacy of a vaccine, or for vaccinating a patient against an infection by a microorganism. The present sequence is that of a PCR primer which was used to amplify and sequence human VH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes an isolated nucleic acid (I) comprising single nucleotide polymorphism (SNP) chosen from: (i) a nucleic acid of Salmo salar SNPs, Oreochromis niloticus SNPs or Atlantic halibut SNPs; and (ii) a nucleic acid having nucleotide sequence that hybridises to (i), or its complement under highly stringent hybridisation conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single nucleotide polymorphism; SNP; fish; Salmo salar; Orechromis niloticus; Atlantic halibut; microsatellite; cod; polymorphic site; seabass; salmonidae; Tilapia; rainbow trout; halibut; detection; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated nucleic acid molecule comprising single nucleotide polymorphism associated with fish, useful for forming PCR primers which are used for detecting single nucleotide polymorphisms in fish nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and VL gene fragments during the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oreochromis niloticus SNP OLA primer SEQ ID NO:302.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 6; SEQ ID NO 302; 233pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         854 AGGAGGAGCTGGTGGAGGCTG 874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 AGGTGCAGCTGGTGAGTCTG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-JAN-2003; 2003WO-IB000112.
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16-AUG-2002; 2002US-0404200P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD19667 standard; DNA; 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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CC Also described: (1) an isolated oligonucleotide (II) comprising at least CC inlotticus SNPs, 0. Inlotticus microsatellites; Atlantic hallbut SNPs, 0. Or plotticus SNPs, 0. Inlotticus microsatellites; Atlantic hallbut SNPs, Cod polymorphic sites and seabass polymorphic sites, or their complement; (2) or amplifying a nucleotide sequence chosen from S. salar SNPs and, 0. Inlotticus SNPs, 0. Inlotticus microsatellites, Atlantic hallbut SNPs, cod polymorphic sites and seabass polymorphic sites; and determining (MI) the corrigin of fish sample comprising providing a parentage genotype database comprising a collection of candidate parent genotype database comprising a collection of candidate parent genotype database, where a match between candidate parent genotype to the parentage genotype database, where a match between the sample genotype to the parentage genotype database, where a match between the sample genotype to the parentage genotype database, where a match between the sample genotype and one of the condidate parent genotype in the sample such as family salmonidae, S. salar, Tilapia, O. niloticus, crainbow trout, halibut, seabass and Atlantic cod. (II) is useful for detecting nucleic acid molecule comprising SNP in a sample, which crainvolus nucleic acid molecule comprising snotleic acids with one or more (II) derived from nucleotide sequence of S. salar SNPs and O. niloticus SNPs, and identifying nucleic acid molecule comprising a polymorphic acids with one or more (II) which is derived from O. niloticus SNPs, Atlantic halibut SNPs, cod polymorphic acides with one or more (II) which is derived from O. niloticus SNPs, Atlantic hilbut SNPs, cod polymorphic acides with one or more (II) which is derived from O. niloticus SNPs, Atlantic hilbut SNPs, cod polymorphic acides with one or more (III) which is derived from O. niloticus sites, and identifying nucleic acide molecule comprising a microsatellite sequence in sample. Comprising a microsatellite sequence in sample. The present sequence is sequence of S. sup
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polynucleotide useful for detecting single nucleotide polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; single nucleotide polymorphism; microarray; side effect; 88; primer; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphism detection primer, SEQ ID No 1842.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 BP; 7 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                730 AAGGAGCTAGAGGTTCTCTCC 750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 AAGGAGATAGAGATGCTCTCC 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF88259 standard; DNA; 23 BP
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-820454/77.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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from a human gene having any one of 335 fully defined sequences as given in specification, or a sequence having a base substitution. The invention further relates to: an oligonucleotide containing single nucleotide polymorphisms; a PCR primer set chosen from the combination of two DNA fragments from any one of 1220 fully defined sequences as given in specification; a labelling probe containing the SNP containing oligo, and a microarray equipped with the SNP containing oligo. The isolated human gene of the invention is useful for detecting the single nucleotide polymorphisms in human gene. The isolated human gene is also useful for disease and determination of side effect to a medical agent. The isolated human gene is also effective in detecting single nucleotide polymorphisms in a human gene. This polymorphism in a human gene. This polymorphism is a human gene. This polymorphism con of the PCR primers used in the single nucleotide polymorphism
The invention relates to a novel polynucleotide isolated and purified
8X868888888888888888888
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ö 0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03; 3; Indels Sequence 23 BP; 10 A; 9 C; 2 G; 2 T; 0 U; 0 Other; 0; Mismatches 2331 GTGCGTGTGTGTGTGTGTG 2351 23 grácarargrárcrárgrára 3 Best Local Similarity 85.7 Matches 18, Conservative Query Match ò

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Gaps

ABX90512 standard; DNA; 23 BP.

ABX90512;

Human VEGFR-2 RT-PCR primer #2.

(first entry)

01-MAY-2003

Antisense; ss; PCR; VEGF; vascular endothelial growth factor; human; cancer; angiogenesis; neoplastic proliferation; cellular proliferation; primer; RT-PCR; reverse transcriptase PCR.

Homo sapiens.

US2002165174-A1.

07-NOV-2002.

13-MAR-2001; 2001US-00805761.

30-JAN-1998; 98US-00016541. 19-JAN-2000; 2000US-00487023. 19-JAN-2001; 2001WO-US000019. 31-JAN-1997; 97US-0037004P.

(GILL/) GILL P S. (MASO/) MASOOD R.

Gill PS, Masood R;

WPI; 2003-255224/25.

New composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor, useful for preparing a composition Example 12; Page 19; 54pp; English vascular endothelia for treating cancer

The invention relates to a composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor (VEGF). The antisense oligonucleotide is useful for preparing a composition treating cancer, neoplastic proliferation, abnormal cellular proliferation and preventing analogenesis. The present sequence is a reverse transcriptase (RT)-PCR primer for a VEGF or related gene, used to

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The invention relates to assessing a culture of undifferentiated primate pluripotent stem cells (PPS, e.g. embryonic stem cells), involving capteression of markers (MR1) e.g. Zic family member 3 (ZIC3), as given in specification, other than human telomerase reverse transcriptase (MR2) and as excipto or podocalyxin-like protein and hrERT and/or Oct3/4 or second marker chosen from (MR2). Also included are maintaining (M2) or second markers (MR2) at a higher level, FOXOLA, ZIC3, hypotherical protein FLJ20582, Forkhead box H1 (FOXH), Zinc finger protein, Hsal2, KRAB-zinc finger protein SZF1-1 or zinc finger protein Gerebellum (ZIC2, or any other marker (WR4) chosen from PHD protein Jade-1, (Jade-1), causing pPS cells (ZNF300), etc., as given in the specification), causing pPS cells (ZNF300), etc., as given in the tissue type by causing them to express one of the markers chosen from
                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Assessing culture of undifferentiated primate pluripotent stem cells by detecting expression of markers e.g., Zic family member 3, other than human telomerase reverse transcriptase/octamer binding transcription
clone the coding region for expression in tumour cell lines. The cell
                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                              Human; ss; PCR; embryonic stem cell; pluripotent stem cell; abnormal cell growth; malignancy; differentiation; probe; RTQ-PCR; realtime quantitative PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mandalam R;
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                  lines were used to test prospective antisense oligonucleotides
                                                                              Length 23;
                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Irving JM,
                                              Sequence 23 BP; 4 A; 7 C; 7 G; 5 T; 0 U; 0 Other;
                                                                         Score 16.2; DB 1;
Pred. No. 1.4e+03;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; SEQ ID NO 41; 106pp; English.
                                                                                                                                           1950 GATCATGCGGGAGTGCTGGCA 1970
                                                                                                                                                            21 GACCATGCTGGACTGCTGGCA
                                                                                                                                                                                                                                                                                                                                                    RTQ PCR probe for Human SLC6A8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brandenberger R,
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                                                                             ttch 0.4%;
sal Similarity 85.7%;
18; Conservative (
                                                                                                                                                                                                                                                      ADI82211 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                     (first entry)
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BRANDENBERGER
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IRVING J M.
MANDALAM R.
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                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shelton D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mok M,
                                                                                                                                                                                                                       RESULT 957
                                                                                                            Matches
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Core promoter element binding protein (COPEB), etc., as given in the cope promoter element binding protein (COPEB), etc., as given in the cope promoter element binding protein (COPEB), etc., as given in the cope appearation), maintaining PBS cells in a pluripotent state (involves culturing PBS cells or their progeny in the presence of a normally differentiation of human embryonic stem (hES) cells, chosen from children per content of the specification), causing pBS cells to differentiate from commally secreted protein that is encoded by a gene that up-regulated upon the specification), causing pBS cells to differentiate (involves culturing pPS cells or their progeny in the presence of a normally secreted protein that is encoded by a gene that up-regulated upon to protein 1 gene, KlaAd085 protein gene, Fax 19) gene, etc., as given in the specification), causing an encoding sequence to be preferentially expressed in undifferentiated cells, causing an encoding sequence to be preferentially expressed in differentiated cells, cope proliferate without differentiation is dentifying genes that are up or secreted during (M4) differentiated cells from less differentiated cells from less differentiated cells from serving (M4) differentiated cells from less differentiated cells from less differentiated cells from less differentiated cells from a sessing culture of ppg-lation, identifying genes that are up or down regulated during differentiation, identifying genes that are up or come of proliferate without differentiated primate pluripotent seem cells and sassessing culture of ppg-lation of ppS cells from human patient of ppg-lation and assessing culture of undifferentiated primate pluripotent seem cells and from a human patient suspected of having a clinical condition related to abnormal cell growth. The method further involves determining whether the patient has a malignancy from the marker expression. The present sequence is an RTO-PCR probe (realtime quantitative).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Transgenic bovine foetus human heavy chain locus PCR primer #1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03; rive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23 BP; 4 A; 9 C; 6 G; 4 T; 0 U; 0 Other;
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Tomizuka K, Ishida I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAAGGCCTGGCTACGCCTCC 23
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08-NOV-2002; 2002US-0425056P.
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Kuroiwa Y,
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WPI; 2004-022868/02

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                                                                                                              The present invention relates to a new transgenic ungulate which comprises one or more nucleic acids encoding all or part of a xenogenous immunoglobulin (Ig) gene which undergoes rearrangement and expresses more than one xenogenous Ig protein. The transgenic ungulate is useful in producing large quantities of xenogenous antibodies, e.g., human antibodies. The present sequence is a PCR primer used in the exemplification of the invention
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            New transgenic ungulate comprising one or more nucleic acids encoding or part of a xenogenous immunoglobulin (Ig) gene, useful in producing large quantities of xenogenous antibodies, e.g., human antibodies.
                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ungulate; human; immunoglobulin; Ig; antibody; transgenic; PCR;
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                                                                                                                                                                                                                                                                                        0.4%; Score 16.2; DB 1; Length 23; B5.7%; Pred. No. 1.4e+03;
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                                                                                                                                                                                                                                                        Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                            854 AGGAGGAGCTGGTGGAGGCTG 874
                                                                                                                                                                                                                                                                                                                                                                                  2 AGGTGCAGCTGGTGTGTG 22
                                                                                  Example 1; Page 74; Opp; English.
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Tomizuka K, Ishida I
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08-NOV-2002; 2002US-0425056P.
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACH01329;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 959
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X444X8X0000000X8
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bovine; non-naturally occurring mutation; mutation; prion protein; mutant prion protein; antibody; immunoglobulin; Ig; agricultural; prion-related disease; bovine spongiform encephalopathy; mad cow disease;

BSE; PCR; primer; ss

sapiens.

Ношо

Synthetic.

WO2004044156-A2.

27-MAY-2004

Human Cmu and VH rearrangment related PCR primer SEQ ID NO:11.

12-AUG-2004 (first entry)

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                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Transgenic bovine locus human heavy chain locus PCR primer #9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kasinathan P, Goldsby RA;
                    Length 23;
                                                                                      3; Indels
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                        Score 16.2; DB 1;
Pred. No. 1.4e+03;
0.4%; Sco...
85.7%; Pred. No. 1...
0; Mismatches
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Tomizuka K, Ishida I;
                                                                                                                                                           854 AGGAGGAGCTGGTGGAGGCTG 874
                                                                                                                                                                                                 2 AGGIGCAGCIGGIGGAGICIG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 79; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                           ACH01336 standard; DNA; 23 BP.
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08-NOV-2002; 2002US-0425056P
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                    Query Match
Best Local Similarity 85.73
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-022868/02.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003097812-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Robl JM, Co
Kuroiwa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACH01336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                         ACHO1336

ACHO1336

AC ACHO136

AC ACHO136
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Collas P, Sullivan E, Kasinathan P;

08-NOV-2002; 2002US-0425056P. 26-SEP-2003; 2003US-0506901P. 10-NOV-2003; 2003WO-US035720.

(HEMA-) HEMATECH LLC. (KIRI) KIRIN BREWERY KK.

Robl J, Kuroiwa Y, Co Tomizuka K, Ishida I;

WPI; 2004-420302/39

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The present invention describes a bovine (I) or a bovine cell (II), comprising a non-naturally occurring mutation in one or both alleles of comprising a non-naturally occurring mutation in one or both alleles of an endogenous prion nucleic acid. Also described: (I) a hybriddoma (III) comprising (II); and (4) a nucleic acid (IV) comprising (III); casette which includes in 5'-3' order a first region of homology having cubstantial sequence identity to a first region of homology having substantial sequence identity to a second region of homology having substantial sequence identity to a second region of homology having substantial sequence identity to a second cregion of prion nucleic acid, first region of homology is longer than second region of homology, where the cassette is capable of integrating into an endogenous prion nucleic acid of cell. (I) is useful for crecovering antibodies, which involves administering one or more antigens of interest to (I) comprising nucleic acid encoding a xenogenous antibodies from (I). The antibodies are monoclonal crecovering xenogenous antibodies from (I). The antibodies are monoclonal crecovering xenogenous antibodies from (I). The antibodies and source of antigen. (I) is useful as source of agricultural products and as human antibodies for pharmaceutical use. (I) is a safe source of agricultural products and as human antibodies for pharmaceutical use. (I) is a safe source of agricultural products and as human antibodies con human antibodies and as human anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel transgenic bovine or bovine cell comprising non-naturally occurring mutation in one or both alleles of endogenous prion nucleic acid, useful as safe source of agricultural products and human antibodies for pharmaceutical use.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (I) has resistance to prion-related disease such as bovine spongiform encephalopathy (mad cow disease). The present sequence represents a PCR primer used in the introduction and rearrangement of HAC, which is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
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Pred. No. 1.4e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23 BP; (3, A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          n an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 11; 307pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           854 AGGAGGTGGTGGAGGCTG 874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 Acerecaecreerecaerere 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%;
Local Similarity 85.7%;
les 18; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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854 AGGAGGAGCTGGTGGAGGCTG 874

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AGGTGCAGCTGGTGGAGTCTG 22

BP

ADO36078 standard; DNA; 23

RESULT 961

AD036078;

ADO36078
ID ADO3
XX
AC ADO3

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bovine; non-naturally occurring mutation; mutation; prion protein; mutant prion protein; antibody; immunoglobulin; Ig; agricultural; prion-related disease; bovine spongiform encephalopathy; mad cow disease;
                                                                                                                                                                                                                                                                                                                                       Novel transgenic bovine or bovine cell comprising non-naturally occurring mutation in one or both alleles of endogenous prion nucleic acid, useful as safe source of agricultural products and human antibodies for
                                                                           Human heavy chain rearrangment related PCR primer SEQ ID NO:17.
                                                                                                                                                                                                                                                                                          Collas P, Sullivan E, Kasinathan P;
                                                                                                                                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 17; 307pp; English,
                   ADO36084 standard; DNA; 23 BP
                                                                                                                                                                                                                                 08-NOV-2002; 2002US-0425056P-26-SEP-2003; 2003US-0506901P-
                                                                                                                                                                                                              10-NOV-2003; 2003WO-US035720
                                                         (first entry)
                                                                                                                                                                                                                                                             (HEMA-) HEMATECH LLC.
(KIRI ) KIRIN BREWERY KK.
                                                                                                                                                                                                                                                                                                   romizuka K, Ishida I;
                                                                                                                          BSE; PCR; primer; ss.
                                                                                                                                                                                                                                                                                         Kuroiwa Y,
                                                                                                                                                                                                                                                                                                                      WPI; 2004-420302/39
                                                                                                                                                                                                                                                                                                                                                                     pharmaceutical use.
                                                                                                                                                                         WO2004044156-A2
                                                                                                                                              Homo sapiens.
                                                         12-AUG-2004
                                                                                                                                                                                           27-MAY-2004
                                                                                                                                                       Synthetic.
                                      ADO36084;
                                                                                                                                                                                                                                                                                          Robl J,
RESULT 962
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The present invention describes a bovine (I) or a bovine cell (II), comprising a non-naturally occurring mutation in one or both alleles of an endogenous prion nucleic acid. Also described: (I) a hybridoma (III) formed from the fusion of (II) with myeloma cell; (2) producing (M1) (I); (3) producing (M2) (II); and (4) a mucleic acid (IV) comprising a cassette which includes in 5'-3' order a first region of homology having substantial sequence identity to a first region of homology having substantial sequence identity to a first region of homology having substantial sequence identity to a second region of homology having substantial sequence identity to a second region of homology where the cassette is capable of integrating region of homology, where the cassette is capable of integrating into an endogenous prion nucleic acid of cell. (I) is useful for producing antibodies, which involves administering one or more antigens of interest to (I) comprising nucleic acid encoding a xenogenous antibody gene locus, the nucleic acid segments in gene locus undergo rearrangement resolving in the production of antibodies specific for the antigen and recovering xenogenous antibodies from (I). The antibodies are monoclonal or polyclonal antibodies. The antibodies are recovered from the serum or milk of bovine. The immunoglobuling (Igs) are directed against a desired milk of bovine. The immunoglobulins (Igs) are directed against a desired antigen. (I) is useful as source of agricultural products and source of human antibodies for pharmaceutical use. (I) is a safe source of agricultural products and as human antibodies for pharmaceutical use, as (I) has resistance to prion-related disease such as bovine spongiform encephalopathy (mad cow disease). The present sequence represents a PCR primer used in the introduction and rearrangement of HAC, which is used in an example from the present invention

DB 1; Length 23; Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other; 0.4%; Score 16.2; Query Match

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                   Gaps
                   ;
Pred. No. 1.4e+03;
                  0; Mismatches
                                     874
                                     854 AGGAGGAGCTGGTGGAGGCTG
                                                  AGGIGCAGCIGGIGGAGICIG
         Similarity 85.7%;
18; Conservative (
          Best Local Similarity
                  Matches
                                      ਨੇ
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bovine; non-naturally occurring mutation; mutation; prion protein; mutant prion protein; antibody; immunoglobulin; Ig; agricultural; prion-related disease; bovine spongiform encephalopathy; mad cow disease; Human heavy chain rearrangment related PCR primer SEQ ID NO:24. ADO36091 standard; DNA; 23 BP (first entry) 12-AUG-2004 AD036091; ADO36091 셤

BSE; PCR; primer; ss WO2004044156-A2 sapiens. 27-MAY-2004 Synthetic. Homo

10-NOV-2003; 2003WO-US035720

08-NOV-2002; 2002US-0425056P. 26-SEP-2003; 2003US-0506901P.

Kasinathan P; Sullivan E, Collas P, (HEMA-) HEMATECH LLC. (KIRI) KIRIN BREWERY KK. Robl J, Kuroiwa Y, Co Tomizuka K, Ishida I; Robl J,

occurring

WPI; 2004-420302/39.

occurring Novel transgenic bovine or bovine cell comprising non-naturally occurring mutation in one or both alleles of endogenous prion nucleic acid, useful as safe source of agricultural products and human antibodies for pharmaceutical use.

Example 2; SEQ ID NO 24; 307pp; English

comprising a non-naturally occurring mutation in one or both alleles of an endogenous prion nucleic acid. Also described: (1) a hybridoma (III) formed from the fusion of (II) with myeloma cell; (2) producing (MI) (1); (3) producing (M2) (II); and (4) a nucleic acid (IV) comprising a cassette which includes in 5'-3' order a first region of homology having substantial sequence identity to a first region of an endogenous prion nucleic acid of a bovine cell, positive selection marker and second region of prion nucleic acid, first region of homology is longer than second region of homology, where the casette is capable of integrating second region of homology, where the casette is capable of integrating into an endogenous prion nucleic acid of cell. (1) is useful for producing antibodies, which involves administering one or more antibods of interest to (1) comprising nucleic acid encoding a xenogenous antibody gene locus, the nucleic acid segments in gene locus undergo rearrangement resulting in the production of antibodies specific for the antigen and recovering in the production of antibodies specific for the antigen and or polyclonal antibodies. The antibodies are recovered from the serum or milk of bovine. The immunoglobulins (Igs) are directed against a desired antigen. (I) is useful as source of agricultural products and source of agricultural products and source of agricultural products and as human antibodies for pharmaceutical use. (I) is a safe source of agricultural products and as human antibodies for pharmaceutical use. (I) has resistance to prion-related disease such as bovine spongiform The present invention describes a bovine (I) or a bovine cell (II)

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RNA-protein fusion; protein library; protein isolation; DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Producing protein or DNA libraries which are useful for improving existing proteins, by in vitro translating protein coding sequences to produce RNA-protein fusions and incubating these protein fusions under high salt conditions.
                                                                                                     RNA-protein fusion oligonucleotide 40-P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 44; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                  01-FEB-2000; 2000WO-US002589.
                          AAA94320 standard; DNA; 39
                                                                                                                                                                                                                                                                                                                                                                                                                    GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-533022/48.
                                                                                                                                             gene cloning; ss.
                                                                                                                                                                                                                                                                                                               WO200047775-A1
                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                                                                                                                           09-FEB-1999;
                                                                              11-JAN-2001
                                                                                                                                                                                                                                                                                                                                        17-AUG-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Szostak JW,
                                                                                                                                                                       Synthetic.
                                                    AAA94320;
                                                                                                                                                                                                           misc_RNA
                AAA94320/
                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to novel antibody sequences, which acts against lesioned tissue. Also claimed is a method (MI) for isolating polynucleotide encoding the antibodies, which involves (a) isolating B cells that is infiltrated into lesioned tissue, and (b) acquiring polynucleotide that encodes an antibody from the isolated B cells. The antibodies are useful for treating cancer lesions, arteriosclerosis, inflammatory disease or autoimmune disease. The present sequence was used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolating polymucleotide that encodes antibody which acts against lesioned tissue, involves isolating B cells that is infiltrated into lesioned tissue, and acquiring polymucleotide that encodes antibody from
encephalopathy (mad cow disease). The present sequence represents a PCR primer used in the introduction and rearrangement of HAC, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                         Cytostatic; Antiarteriosclerotic; Antiinflammatory; Immunosuppressive; human; antibody; cancer lesion; arteriosclerosis; inflammatory disease; autoimmune disease; cancer; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tsunoda H;
                                                                                                         Gaps
                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 16.2; DB 1; Length 23; 85.7%; Pred. No. 1.4e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matsubara K,
                                                                              Length 23;
                                                                                                       3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                                     Sequence 23 BP; 3 A; 4 C; 11 G; 5 T; 0 U; 0 Other;
                                                                            0.4%; Score 16.2; DB 1;
85.7%; Pred. No. 1.4e+03;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fujii E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 104; 200pp; Japanese.
                                                                                                                                                                                                                                                                                                       Antibody related primer, SEQ ID 104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yoshida K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHUS ) CHUGAI SEIYAKU KK.
(PHAR-) PHARMALOGICALS RES PTE LTD.
                                                                                                                                  854 AGGAGGAGCTGGTGGAGGCTG 874
                                                                                                                                                          2 Agerecaecreerecaerere 22
                                                                                                                                                                                                                          ADP03731 standard; DNA; 23 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-NOV-2003; 2003WO-JP014919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-NOV-2002; 2002JP-00339241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to illustrate the invention.
                                                                                                                                                                                                                                                                             26-AUG-2004 (first entry)
                                                                                            Best Local Similarity 85.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Psuchiya M, Suzuki M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-450382/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           isolated B cells.
                                                                                                                                                                                                                                                                                                                                                                                                            WO2004048571-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                     ADP03731;
                                                                                Query Match
                                                                                                                                                                                                   RESULT 964
                                                                                                                                                                                                               ADP0373
                                                                                                                                                                                                                            ន្តដូច្ចប្រ
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Liu R;

Roberts RW,

'note= "attached to puromycin, a peptide acceptor"

/*tag= b /mod_base= OTHER

Location/Qualifiers

(first entry)

/*tag= a /label= RNA

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The present sequence is one of a number of oligonucleotides which were used for the generation of RNA-protein fusions, including fusions having used for the generation of RNA-protein fusions comprise a protein covalently any or epitope tag. The RNA-protein fusions complished by synthesis and in vitro or in situ translation of an mRNA molecule with a peptide and in vitro or in situ translation of an mRNA molecule with a peptide acceptor attached to its 3' end. The RNA-protein fusions are incubated under high salt conditions to produce a protein library. This method is useful for improving or altering existing proteins, as well as for isolating new proteins and nucleic acid or small molecule targets. It may can be used to improve human or humanised simple-chain antibodies for the treatment of a number of diseases. The method is useful for the case and cloning new genes on the basis of protein-protein interactions. Unlike prior art, the new method does not rely on complex, which is very fragile and is therefore of limited use. The method does not rely on topological links between the protein and the nucleic acid so that the information of the protein is retained and can conclete acid in readable, nucleic acid form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.4%; Score 16.2; DB 1; Length 39; Best Local Similarity 72.4%; Pred. No. 2.1e+03; Matches 21; Conservative 0; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 39 BP; 28 A; 4 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3260 GATATTTTATTTGCTTTGTCCTTTTCAG 3288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ò
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Gaps

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854 AGGAGGAGCTGGTGGAGGCTG 874

18; Conservative

Query Match Best Local Similarity Matches 18; Conserv

AGGTGCAGCTGGTGGAGTCTG 22

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02-FEB-1993 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MboI DNA fragments of between 250 and 500 by with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites is estimated at >100,000. The sequence information for ca. 230 such bovine microsatellites is summarised in the sequence information of commetream of the microsatellite sequence were used to generate the carpired PCR primers for in vitro amplification of the corresp.

The sequence were used to generate the microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                          - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                 PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 9.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 16 BP; 0 A; 0 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                         Microsatellite sequence from clone TGLA158.
 39 GGTTTTTTTTTTTTTTTTTTCAG 11
                                                                                                                                                                                                                                                                                                                                                                                                                              Table 7; Page 227; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic bovine DNA markers - mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2318 TGTGTGTGTGTGTG 2333
                                                        AAQ33743 standard; DNA; 16 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGTGTGTGTGTGTGTG 16
                                                                                                                                                                                                                                                                                                91US-00642342
                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                              Georges M, Massey JM;
                                                                                                        (revised)
                                                                                                                                                                                                                                                                                                                                                                    WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
les 16; Conserv
                                                                                                                                                                                                                                                                                                                       (GENM-) GENMARK
                                                                                                       25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                          WO9213102-A1
                                                                                                                                                                                                                                                                         15-JAN-1992;
                                                                                                                                                                                                                                                                                                15-JAN-1991;
                                                                                                                                                                                                                                                  06-AUG-1992
                                                                                                                                                                                                    Bos taurus.
                                                                                 AAQ33743;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                   RESULT 966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                           셤
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screening a library of bovine MoI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MbOI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for oa. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequence upstream and downstream of the microsatellite sequence were used to generate the microsatellite (using the program Optibication of the corresp. microsatellite (using the program Optibication and individuals, for parentage testing, and in the genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic bovine DNA markers - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequence is that of a bovine microsatellite sequence obtd. by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                           PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 16; 9.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16 BP; 0 A; 0 C; 8 G; 8 T; 0 U; 0 Other;
Microsatellite sequence from clone TGLA160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 16; DB
100.0%; Pred. No. 9.7
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Microsatellite sequence from clone TGLA311.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Table 7; Page 229; 517pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mapping, and selective breeding.
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(first entry)
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Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                  (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                             15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                              15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9213102-A1
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02-FEB-1993
                                                                                                                                                                                                                                         36-AUG-1992
                                                                                                                                   Bos taurus.
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AAQ33749 standard; DNA; 16 BP

RESULT 967 AAQ33749

25-MAR-2003 (revised)

AAQ33749;

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                                                                                                                                                                                                                        The sequence is that of a bovine microsatellite sequence obtd. by
screening a library of bovine MboI DNA fragments of between 250 and 500
by with an (ACJIS and a (TC)15 oligomucleotide probe. One out of 50
clones cross-hybridised. Assuming independent distribution of
microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites
in the bovine genome is estimated at >100,000. The sequence information
cross-fication and indexed herein (see below). The sequence information
cross-fication and indexed herein (see below). The sequences upstream and
downstream of the microsatellite sequence ware used to generate the
required PCR primers for in vitro amplification of the corresp.
microsatellite (using the program OPTIPRIM). The microsatellites may be
used to identify individuals, for parentage testing, and in the genetic
mapping of economic trait loci, or genes involved the determinism of
economically important traits esp. in cattle, to allow selective
breeding. See also AAQ33501-34437. (Updated on 25-WAR-2003 to correct PN'
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                                                                                                                                                              Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding.
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oligonucleoside; ss.
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Pred. No. 9.7e+02;
0; Mismatches 0; Indels
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100.0%; Pred. No. ...
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                                       92WO-US000340
                                                                91US-00642342
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                 Massey JM;
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es 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENT-) GENTA INC.
                                                                                         (GENM-) GENMARK
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                                        15-JAN-1992;
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16-FEB-1995
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               06-AUG-1992
                                                                                                                 Georges M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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AAQ68236/c
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Matches
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                                                                                                                                                                                                                                                                  Two sets of methylphosphonate oligonucleosides ("MP oligomers") and complementary ribooligonucleosides ("RNA oligomers") contg. alternating purines and pyrimidines were examined for their ability to form triple helix complexes. (Set 1:G2019 and R138; Set 2:G2018 and R139; see AAQG8235-38). It was shown that MP oligomers contg. alternating purines and pyrimidines are capable of forming triple stranded complexes with complementary RNA oligomers. (Updated on 25-MAR-2003 to correct PN
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                                                        Detection, recognition, inhibition and alteration of single and double stranded target nucleic acid sequences - by formation of a triple helix structure using 2 oligomers which block translation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence 16 BP; 8 A; 8 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                 Example 2; Page 37; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ68235 standard; DNA; 16 BP
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WPI; 1994-217542/26.
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Best Local Similarity
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16-FEB-1995
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complementary ribooligonucleosides ("RNA oligomers") contg. alternating purines and pyrimidines were examined for their ability to form triple helix complexes. (Set 1:G2019 and R138; Set 2:G2018 and R139 - see AAQ68235-38). It was shown that Williams contg. alternating purines and pyrimidines are capable of forming triple stranded complexes with complementary RNA oligomers. (Updated on 25-WAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                          Detection, recognition, inhibition and alteration of single and double stranded target nucleic acid sequences - by formation of a triple helix structure using 2 oligomers which block translation.
                                                                                  Purine; pyrimidine; methylphosphonate; MP; triple helix; translation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Two sets of methylphosphonate oligonucleosides ("MP oligomers") and
                                                    Purine-pyrimidine contg. methylphosphonate oligonucleoside G2018
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                                                                                                                                                                                                                                           92US-00987746.
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           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                  Arnold LJ, Reynolds MA;
                                                                                                                                                                                                                                                                                                                                WPI; 1994-217542/26.
                                                                                               oligonucleoside; ss
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les 16; Conserv
                                                                                                                                                                                                                                                                       (GENT-) GENTA INC.
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18-JUN-1997
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                                                                                                                            Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detection, recognition, inhibition and alteration of single and double stranded target nucleic acid sequences - by formation of a triple helix structure using 2 oligomers which block translation.
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                                                                    Gaps
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50.0%; Pred. No. 9.7e+02;
iive 8; Mismatches 0; Indels
                                     0.4%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 9.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                    Purine-pyrimidine contg. ribooligonucleoside R139.
             Sequence 16 BP; 0 A; 0 C; 8 G; 8 T; 0 U; 0 Other;
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                                                                                               2335 GTGTGTGTGTGTGT 2350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                  AAQ68238 standard; RNA; 16 BP
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                                                     Local Similarity 100.
es 16; Conservative
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                                                                                                                                                                                                                                                           (revised)
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16-FEB-1995
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                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                               AAQ68238;
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                                         Query Match
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                                                    Gaps
                                                    ö
                      0.4%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 9.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                  Repeat sequence found in ADP/ATP translocase gene.
Sequence 16 BP; 8 A; 8 C; 0 G; 0 T; 0 U; 0 Other;
                                  100.0%; Prec. ...
                                                                              2335 GTGTGTGTGTGTGT 2350
                                                                                                                                                                         AAT66090 standard; DNA; 16 BP.
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                                                                                             16 Grerererererer
                                                     Conservative
                                                                                                                                                                                                                              (revised)
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AAQ68237 standard; DNA; 16 BP

RESULT 972
AAQ68237/c
ID AAQ682
XX
AC AAQ682

AAQ68237

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Gaps

Weber JL;

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The invention relates to a substantially purified or isolated nucleic acid (1) from ryegrass or fescue species including a simple sequence repeat (SSR), having 2 or more tandemly repeated nucleotide core elements 2-6 nucleotides in length. Also included are a nucleic acid primer suitable for amplifying an SSR, identifying (M1) an SSR by preparing a library of ryegrass or fescue genomic DNA enriched for SSRs and identifying clones in the library containing SSRs, a library of ryegrass or fescue genomic DNA enriched for SSRs prepared by the M1, selecting for a gene in grass or cereal breeding by identifying an SSR that is closely associated with the gene such that the SSR and the gene are associated with the gene such that the SSR and the gene are preferentially co-inherited, and selecting for the SSR in the breeding, a
                    method which is not dependent on base sequences, then selection of the statellite sequences from the obtained genomic library of high homogeneity. The method is useful for the isolation of microastellite DNA sequences which can be used as DNA markers. The new method markedly improves the efficiency of isolation of satellite sequences in comparison to prior art methods which are reliant on base sequences. Sequences AA296483-514 represent sequences from Haliotis discus, used in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New simple sequence repeats having 2 or more tandemly repeated nucleotide core elements isolated from ryegrass and fescue, useful for selecting of genes in grass or cereal breeding or profiling grass or cereal species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SSR; ryegrass; fescue; tandem repeat; seed batch purity testing.
sequences from genomic DNA that comprises fragmentation of the DNA by
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                                                                                                                                                                                                                                                             Length 16;
                                                                                                                                                                                                                                                                                                        0; Indels
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UNIV ADELAIDE.
                                                                                                                                                                                                                     Sequence 16 BP; 8 A; 8 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                           0.4%; Score 16; DB 1; L
100.0%; Pred. No. 9.7e+02;
:ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence repeat; plant; ds; profiling; grass profiling;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Simple sequence repeat, SSR, #42.
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                                                                                                                                                                                                                                                                                                                                                      2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAS13770 standard; DNA; 16 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-MAY-2000; 2000AU-00007310.
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                                                                                                                                                                                                                                                                                      Local Similarity 100.
nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones ES;
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                                                                                                                                                                            of the invention
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                                                                                                                                                                                                                                                                  Query Match
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(VICT-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolation of satellite sequences from genomic DNA for use as DNA markers comprises isolating a library with high homogeneity by DNA fragmentation.
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                                                                                                                                                                                                                                             Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n using novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Pred. No. 9.7e+02;
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100.0%; Pred. No. ...
... 0; Mismatches
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                                                                                                                                                                                                                                                                                                                 Example 9; Col 59-60; 186pp; English.
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         94US-00222177.
                                                 89US-00341562,
91US-00754351,
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Best Local Similarity 100.0
Matches 16; Conservative
                                                                                                                     (MARS-) MARSHFIELD CLINIC.
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                                                                                                                                                                                                         WPI; 1997-042299/04
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             04-APR-1994;
                                                    21-APR-1989;
05-SEP-1991;
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AAZ98508;

RESULT 974 AAZ98508,

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method for DNA profiling grass or cereal species varieties by assessing variation between SSR varieties and testing the purity of grass or cereal seed batches by assessing variation within seed batch of an SSR. The SSRs may be used in the selection of genes in grass or cereal breeding, for profiling grass or cereal species varieties, for testing the purity of grass or cereal seed batches, and for DNA profiling to establish the electronic and for DNA profiling to establish the sequence is a ryegrass or fescue SSR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention discloses a method for generating five prime biased tandem tag libraries of cDNAs. The step involves isolating a sample of mRNAs, amplifying the released tags, concatenating the amplified tags to form concatenated tags, amplifying and isolating the concatenated tags. The present sequence is human carboxypeptidase A3 tandem tag DNA
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                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                      0.4%; Score 16; DB 1; Length 16;
100.0%; Pred. No. 9.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                      Sequence 16 BP; 0 A; 1 C; 7 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human carboxypeptidase A3 tandem tag DNA #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure, Page 5; Opp, English.
                                                                                                                                                                                                                                                           2315 GTCTGTGTGTGTGT 2330
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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Best Local Similarity 100.0
Matches 16; Conservative
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HERMIDA L C.
HOPPA N L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS2003190618-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                              AAD63065;
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(HOPP/)
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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers pecifically priming and allowing amplification of each Voeta gene. VertaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases. Concluding autoimmune disease, hypersensitivity diseases, infectious diseases and neoplastic diseases. Hypersensitivity diseases, infectious diseases, arrophic gastritis. Degenerative nervous system diseases include Multiple atrophic gastritis Degenerative nervous system diseases include multiple atrophic gastritis and as contact with allergens that lead to allergies, Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by viruses such as HIV, fungal infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                              degenerative nervous system disease, graft versus host disease, hypersensitivity disease; infectious disease, neoplastic disease; haddison's disease; atophic gastritis, degenerative nervous system disease, multiple sclerosis, Alzheimer's disease, hypersensitivity disease; type I hypersensitivity, Goodpasture's syndrome; type II hypersensitivity; Goodpasture's syndrome; HIV; fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          such as cancer of the brain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       súch as leukaemias, lymphomas and cancers such as cancer of the brair
breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                human; T-cell associated disease; Vbeta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16 BP; 8 A; 1 C; 0 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                      Human Vbeta gene repeat sequence #137.
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95US-00531241
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                                       ADH70347 standard; DNA; 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast cancer; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                          25-MAR-2004
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RESULT 977
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Gaps

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0; Indels

0.4%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 9.7e+02;

100.0%; Prea.

16; Conservative

Matches

Local Similarity

Query Match

2335 GTGTGTGTGTGTGT 2350

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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each 'Deta gene, UperaRNA or cDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases including autoimmune diseases, degenerative nervous system diseases.

The rejection and disease, hypersensitivity diseases, infectious diseases including autoimmune diseases, hypersensitivity diseases, infectious diseases, and neoplastic disease. Hypersensitivity diseases include multiple atrophic gastritis. Degenerative nervous system diseases include multiple strophic gastritis as contact with allergens that lead to I hypersensitivities such as contact with allergens that lead to allergies. Type II hypersensitivities such as those present in allergies a syndrome and Type IV hypersensitivities such as those caused by viruses such as HIV, fungal infections at those caused by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzhaimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; HIV; fungal infection, Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                Gaps
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                 Length 16;
                     DB 1; Le
              Score 16; DB 1; Pred. No. 9.7e 0; Mismatches
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0.4%; Scc.
100.0%; Pre
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95US-00531241.
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Best Local Similarity 100.0
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          breast cancer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HOOD/) HOOD L E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2002150891-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nbeta gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH70350;
                                                                                                                                                                                                                                                                                                                                  RESULT 978
                                                                                                                                                                                                                                                                                                                                                              ADH70350/
XX
AAC
ADH7
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ADH70
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The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using cas template a DNA sample containing at least one segment of (I); and determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or predisposition to a disease, associated with a gene that includes one or core PML. The method is used to identify microsatellite markers, in a include PML. The method is used to identify microsatellite markers, in a disease and for predisposis of such diseases, especially prion diseases of the predisposis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins.

Cometabolic diseases; also to type genes that encode milk proteins.

Cometabolic diseases; also to type genes that encode milk proteins.

Comparising a polymorphic microsatellite locus.
                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
the yeast genus Candida, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain,
                                                                         breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease predisposition, microsatellite marker, prion disease, cystic fibrosis, malignant hyperthermia syndrome, metabolic disease, milk protein, hormone, transcription factor, p77-blue-vector, sheep,
                                                                                                                                                                                             ö
                                                                                                                                                         Score 16; DB 1; Length 16;
Pred. No. 9.7e+02;
                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sheep prion protein microsatellite locus primer #66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polymorphic microsatellite loci; PML;
                                                                                                                 Sequence 16 BP; 7 A; 0 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                 Mismatches
                                                                                                                                             0.4%; Sc.
100.0%; Pred
0; N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Han Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     microsatellite; PCR; primer; ss
                                                                                                                                                                                                                                       2823 TATATATACATATATA 2838
                                                                                                                                                                                                                                                                                                                                                                         ВЪ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                       16 rarararacararara 1
                                                                                                                                                                                                                                                                                                                                                                           ADO81095 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                     16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Geldermann H, Preuss S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYHO-) UNIV HOHENHEIM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-215730/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       length of amplicons
                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DE10236711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               typing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                               ADO81095;
                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                     RESULT 979
                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                         ADO81095/
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Polymorphic bovine DNA markers - used in genetic identification, gene
                                                                                                                                                         PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                               Microsatellite sequence from clone TGLA188.
                                                                                                                                                                                                                                                                                                                                                                                                                          mapping, and selective breeding.
                                       AAQ33783 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                         Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                 (GENM-) GENMARK.
                                                                                                                                                                                                                          WO9213102-A1.
                                                                                                                                                                                                                                                                           15-JAN-1992;
                                                                                                                                                                                                                                                                                                     15-JAN-1991;
                                                                                        25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                   36-AUG-1992.
                                                                                                                                                                                                  Bos taurus.
                                                                AAQ33783;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 982
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                        AAQ33783
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine Mbol DNA fragments of between 250 and 500 by with an (AC115 and a (TC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and Mbol sites, the frequency of (T6)n >9 microsatellites and microsatellites is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellite is summarised in the specification and indexed herein (see below). The sequence information consumerream of the microsatellite sequence were used to generate the countried PCR primers for in vitro amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding.
                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                         PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 16; DB 1; Length 17;
100.0%; Pred. No. 1e+03;
iive 0; Mismatches 0; Indels
                         0.4%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 9.7e+02;
                                                  0; Indels
Sequence 16 BP; 8 A; 8 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 0 A; 0 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                Sequence of a microsatellite from clone TGLA84.
                     Query Match 0.4%; Score 16; DE Best Local Similarity 100.0%; Pred. No. 9.7 Matches 16; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Table 7; Page 396; 517pp; English.
                                                                             2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                      AAQ34164 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                             92WO-US000340
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                                                                                                                                                                                                                         (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1992-284684/34.
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Les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                           WO9213102-A1.
                                                                                                                                                                                                                                                                                                                                                                                                             15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-1991;
                                                                                                                                                                                                                         25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                     06-AUG-1992.
                                                                                                                                                                                                                                                                                                                                  Bos taurus.
                                                                                                                                                                                                AAQ34164;
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Matches
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92WO-US000340.

(first entry)

(revised)

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The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine Mbol DNA fragments of between 250 and 500 by with an (ACL)15 and a (TCL)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and Mbol sites, the frequency of (T6)n >9 microsatellites in the bovine ganome is estimated at >100,000. The sequence information for ca. 230 such bovine microsatellites is summarised in the sequence information for ca. 230 such bovine microsatellites is summarised in the sequence of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp.

The microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 0 A; 0 C; 8 G; 9 T; 0 U; 0 Other;
Fable 7; Page 242; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX56865 standard; DNA; 17 BP
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TGTGTGTGTGTGTG 2333

2318

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This invention describes have a comprising (a) an RhaseH-activating region (R1) of at least 3 consecutive 2' unsubstituted mucleosides region (R2) of at least 3 consecutive 2' unsubstituted mucleosides (connected by charged internucleosides inth at least one chirally selected internucleoside link. The oligomucleosides of the invention have base sequences complementary to that of tearget RNA. The products of the invention are used to inhibit transcription of target RNA in a cell or organism, they are antisense molecules that also activate RNaseH.

Particularly the oligomucleosides are used in diagnosis and treatment of disease associated with endogenous or foreign gene expression. Use against human papilloma virus is exemplified. The modified internucleoside links improve target specificity, potency and binding internucleoside links improve target specificity, potency and binding resistant to nuclease and so have better in-vivo lifetimes. The chirally resistant to nuclease and so have better in-vivo lifetimes. The chirally elected linkages in R2 allow control of binding affinity. ARASE6855-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 X56881 represent oligonucleosides used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
       Oligonucleoside; RNaseH-mediated cleavage; target; RNaseH; inhibitor; internucleoside linkage; antisense; diagnosis; treatment; disease; binding affinity; Ka; nuclease resistance; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                              New oligo:nucleotide(s) causing cleavage of target RNA - with RNaseH activated segment having charged inter-nucleoside links and second segment with chirally selected links.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleosides that causes RNaseH-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 16; DB 1; Length 17;
100.0%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Repeat sequence found in the angiogenin gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                           Giachetti C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure, Page 137; 165pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2335 GTGTGTGTGTGT 2350
                                                                                                                                                                                                                              93US-00154013.
93US-00154014.
94US-00233778.
94US-00238177.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                            Reynolds MA,
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-254769/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                            GENT- GENTA INC.
                                                                                                                                                                                                                                16-NOV-1993;
16-NOV-1993;
26-APR-1994;
                                                                                                                        WO9513834-A1
                                                                                                                                                                                               16-NOV-1994;
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18-JUN-1997
                                                                                                                                                                                                                                                                                       04-MAY-1994;
                                                                                                                                                             26-MAY-1995
                                                                                                                                                                                                                                                                                                                                                              Arnold LJ,
                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT66099;
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The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. The sequences AAT66084-T66107 represent repeat sequences of low informativeness found in specific human genes. This repeat sequence is found in the angiogenin gene located at chromosomal position 14q11-q13. The sequence is amplified by primers AAT66100-1. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            internucleosidyl linkage; chirality; hybridization; racemic;
                                                                                                                                                                                                                                                         Detection of polymorphic genetic markers of the form (dC-dA) \, n \, (dG-dT) \, n using novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 16; DB 1; Length 17;
100.0%; Pred. No. 1e+03;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 9 A; 8 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                        Example 9; Col 61-62; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Methylphosphonate oligomer 2517-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2318 TGTGTGTGTGTGTG 2333
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hybridisation; chromosome; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-00885126.
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91US-00754351.
                                                                                                            94US-00222177.
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                                                                                                                                                                               (MARS-) MARSHFIELD CLINIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              binding affinity; ss
                                                                                                                                                                                                                                    WPI; 1997-042299/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENT-) GENTA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schwartz DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          L6-NOV-1993;
                             Homo sapiens
                                                                                                             04-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phosphonate
                                                                                                                                       21-APR-1989;
05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5955597-A
                                                       US5582979-A
                                                                                   10-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX91062;
                                                                                                                                                                                                            Weber JL;
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 984
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The invention provides methods for preparing oligomers having phosphonate internucleosidyl linkages of a preselected chirality which hybridize to a target RNA sequence. The method of making comprises: (a) synthesizing a nucleoside dimer, trimer, or terramer with racemic internucleosidyl phosphonate linkages; (b) purifying the racemic nucleoside to a chirally pure nucleoside; and (c) sequentially linking at least 2 of the chirally pure nucleosides to form a synthetic oligomer that is enriched for phosphonate internucleosidyl linkages of a preselected chirality and is complementary to an RNA target sequence. The methods are useful for providing chirally enriched synthetic oligomers RP chirally enriched synthetic oligomers RP compared to oligomers have enhanced binding affinities for RNA compared to oligomers with racemic all methylphosphonate internucleosidyl linkages.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences AAX91054-75 represent oligomers chemically synthesised using
                                                                             Oligomers made using chirally pure nucleoside dimers, trimers, or tetramers with enhanced binding affinities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 1 A; 0 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                            Example 19; Col 39-40; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the method of the invention
                      WPI; 1999-539600/45.
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Score 16; DB 1; Length 17; Pred. No. 1e+03; 0; Mismatches 0; Indels 0.4%; Scc... 100.0%; Pred. No. ... 2335 GTGTGTGTGTGTGT 2350 Local Similarity 100. Les 16; Conservative Query Match Matches

drererererer 16 ò 셤

AAD17599 standard; DNA; 17 BP AAD17599; RESULT 985 AAD17599,

(first entry) 10-DEC-2001 5' variation generator oligonucleotide PCR primer #14.

Genomic DNA analysis; 5' variation generator; 3' fragment generator; endangered animal identification; PCR primer; 8s.

Unidentified

BP1130114-A1

05-SEP-2001,

03-MAR-2000; 2000EP-00200757

03-MAR-2000; 2000EP-00200757

(VHAE-) VAN HAERINGEN LAB BV.

Van Haringen H, Van Haringen WA;

WPI; 2001-572636/65.

Analyzing genomic DNA in a sample, useful for analyzing genes of organisms (e.g. a species or individual) or identifying endangered animals or plants, by using oligonucleotide primers comprising universal variable fragments.

Example 1; Page 6; 23pp; English.

The patent discloses a method and associated kit for analysing genomic DNA in a sample. The method comprises conducting a nucleic acid amplification on the genomic DNA in the sample using both first and

second oligonucleotide primer to produce DNA fragments based on repeat sequences on at least one end of the genomic DNA. The first primer is a 5' variation generator including a repeat sequence and at least one nonrepeat nucleotide. The second oligonucleotide primer is a 3' fragment generator starting within such a genetic distance that amplification of the genomic DNA can be performed and preferably includes inosine. The method is useful for the genetic analysis of an individual organism, particularly of a species or individual. It is also useful for the rapid and straight forward identification of endangered animals or plants. The present DNA sequence is a 5' variation generator oligonucleotide PCR primer 888888888888888

Sequence 17 BP; 8 A; 8 C; 0 G; 1 T; 0 U; 0 Other;

Gaps ö Length 17; 0: Indels Score 16; DB 1; Pred. No. 1e+03; Mismatches .; 0 2318 TGTGTGTGTGTGTG 2333 0.4%; 8 16; Conservative Query Match Best Local Similarity Matches ठ

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17 rerererererere 2

BP. AAD17594 standard; DNA; 17 AAD17594

RESULT 986

AAD17594;

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Gaps

; 0

(first entry) 10-DEC-2001 5' variation generator oligonucleotide PCR primer #9.

Genomic DNA analysis; 5' variation generator; 3' fragment generator; endangered animal identification; PCR primer; ss.

Unidentified

EP1130114-A1.

05-SEP-2001

03-MAR-2000; 2000EP-00200757.

03-MAR-2000; 2000EP-00200757

(VHAE-) VAN HAERINGEN LAB BV

/an Haringen H, Van Haringen WA;

WPI; 2001-572636/65.

Analyzing genomic DNA in a sample, useful for analyzing genes of organisms (e.g. a species or individual) or identifying endangered animals or plants, by using oligonucleotide primers comprising universal variable fragments.

Example 1; Page 6; 23pp; English.

The patent discloses a method and associated kit for analysing genomic DNA in a sample. The method comprises conducting a nucleic acid amplification on the genomic DNA in the sample unaing both first and second oligonucleotide primer to produce DNA fragments based on repeat sequences on at least one end of the genomic DNA. The first primer is a 5 variation generator including a repeat sequence and at least one nongenerator starting within such a genetic distance that amplification of the genomic DNA can be performed and preferably includes inosine. The method is useful for the genetic analysis of an individual organism, particularly of a species or individual. It is also useful for the rapid and straight forward identification of endangered animals or plants. The present DNA sequence is a 5 variation generator oligonucleotide PCR (first entry)

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Analyzing genomic DNA in a sample, useful for analyzing genes of organisms (e.g. a species or individual) or identifying endangered animals or plants, by using oligonucleotide primers comprising universal
                                                                                                                                                                                                                                                                       Genomic DNA analysis; 5' variation generator; 3' fragment generator; endangered animal identification; PCR primer; ss.
                                                                                                                                                                                                                        5' variation generator oligonucleotide PCR primer #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Van Haringen H, Van Haringen WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 6; 23pp; English.
                                                                       AAD17595 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAR-2000; 2000EP-00200757.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-MAR-2000; 2000EP-00200757.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (VHAE-) VAN HAERINGEN LAB BV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-572636/65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            variable fragments.
                                                                                                                                                                                                                                                                                                                                                                                                        EP1130114-A1.
                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-SEP-2001.
                                                                                                                                                                        10-DEC-2001
                                                                                                                      AAD17595;
                                            AAD17599
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Analyzing genomic DNA in a sample, useful for analyzing genes of organisms (e.g. a species or individual) or identifying endangered animals or plants, by using oligonucleotide primers comprising universal
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                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genomic DNA analysis; 5' variation generator; 3' fragment generator; endangered animal identification; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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                                                                    Score 16; DB 1; Length 17;
Pred. No. 1e+03;
0; Mismatches 0; Indels

    variation generator oligonucleotide PCR primer #12.

                     Sequence 17 BP; 0 A; 0 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 8 A; 9 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                       100.0%; Prec. N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Van Haringen WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 6; 23pp; English.
                                                                                                                                                                           2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                         AAD17597 standard; DNA; 17 BP.
                                                                                                                                                                                                                          2 rerererererere 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (VHAE-) VAN HAERINGEN LAB
                                                                                              Local Similarity 100.
Les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-572636/65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              variable fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Van Haringen H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EP1130114-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                           AAD17597;
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                                                                       Query Match
                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                    RESULT 98
AAD17597(
IN AAD17597
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The patent discloses a method and associated kit for analysing genomic DNA in a sample. The method comprises conducting a nucleic acid amplification on the genomic DNA in the sample using both first and second oligonucleotide primer to produce DNA fragments based on repeat sequences on at least one end of the genomic DNA. The first primer is a 5' variation generator including a repeat sequence and at least one nonrepeat nucleotide. The second oligonucleotide primer is a 3' fragment generator starting within such a genetic distance that amplification of the genomic DNA can be performed and preferably includes inosine. The method is useful for the genetic analysis of an individual organism, particularly of a species or individual. It is also useful for the rapid and straight forward identification of endangered animals or plants. The
                                                                                                                                                                                                                                                                                                                                          and straight forward identification of endangered animals or plants. present DNA sequence is a 5' variation generator oligonucleotide PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1e+03; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 1 A; 0 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human Vbeta gene repeat sequence #153.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2318 TGTGTGTGTGTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similario,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH70363;
                                                                                                                                                                                                                                                                                                                                                                                                             primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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2318 TGTGTGTGTGTGTG 2333

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17 rererererereres 2

ADH70776;

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degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity disease; type II hypersensitivity; doodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; trype IV hypersensitivity; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
Vbeta; autoimmune disease;
 associated disease;
                                                                                                                                                                                              99US-00263959.
                                                                                                                                                                                                                  94US-00309335.
95US-00531241.
                                                                                                                                                                                                                                                                                                    WPI; 2004-059052/06.
                                                                                                                                                                                                                                                                               Hood LE, Rowen L;
                                                                                                                oreast cancer; ds.
                                                                                                                                                                                                                                                (HOOD/) HOOD L E.
                                                                                                                                                                                                                                                            (ROWE/) ROWEN L.
                                                                                                                                                       JS2002150891-A1
                                                                                                                                  Homo sapiens.
                                                                                                                                                                                           05-MAR-1999;
                                                                                                                                                                                                                  19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                           17-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                       Vbeta gene
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Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a Disclosure; SEQ ID NO 557; 164pp; English.

including autoimmune diseases, degenerative nervous system diseases, graft versus host disease, hypersensitivity diseases, infectious diseases and neoplastic diseases. Autoimmune diseases include Addison's disease, atrophic gastritis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple allergies, Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those present in mifested in leprosy. Infectious diseases include viral infections caused by viruses such as HIV, fungal infections such as those caused by viruses caused by arasitic infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases such as cancer of the brain, such as leukaemias, lymphomas and cancers such as cancer of the brain breast. The present sequence represents a Vbeta gene repeat sequence. The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, VbecaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases

Sequence 17 BP; 0 A; 0 C; 8 G; 9 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1e+03; cive 0; Mismatches 0; Indels Query Match Best Local Similarity 100.0

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2318 TGTGTGTGTGTGTG 2333 TGTGTGTGTGTGTG 16

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RESULT 990 ADH70776 ID ADH7077

ADH70776 standard; DNA; 17 BP.

Alzheimer's disease; hypersensitivity disease; type I hypersensitivity, allergy; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; type IV, fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; defenserative nervous system disease; multiple sclerosis; human; T-cell associated disease; Vbeta; autoimmune disease; Human Vbeta gene repeat sequence #566. 25-MAR-2004 (first entry)

Homo sapiens.

breast cancer; ds

US2002150891-A1.

17-OCT-2002.

99US-00263959 05-MAR-1999; 94US-00309335. 95US-00531241. 19-SEP-1994; 19-SEP-1995;

(HOOD/) HOOD L E. (ROWE/) ROWEN L.

Rowen L; HOOD LE,

WPI; 2004-059052/06.

Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a Vbeta gene

Disclosure; SEQ ID NO 970; 164pp; English.

The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, vbetaRNA or cDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases including aucoimmune diseases, degenerative nervous system diseases. Crejection and diseases. Autoimmune diseases include Addison's disease, and neoplastic diseases. Autoimmune diseases include Addison's disease. Cretrophic gastritis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those caused by truses such as HIV, fungal infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes. the brain, breast. The present sequence represents a Vbeta gene repeat sequence. lymphomas and cancers such as cancer of eukaemias, such as

Sequence 17 BP; 0 A; 0 C; 8 G; 9 T; 0 U; 0 Other;

Gaps ; 0.4%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1e+03; tive 0; Mismatches 0; Indels Ouery Match 0.4%. Best Local Similarity 100.0° Matches 16; Conservative

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2318 TGTGTGTGTGTGTG 2333

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Gaps

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ADH70511 standard; DNA; 17
WPI; 2004-059052/06
                                                                                                                   Hood LE, Rowen L;
                                                                                                          HOOD L E.
ROWEN L.
                                                                                 US2002150891-A1
                                                                      breast cancer;
                                                                                                          (ROWE/) ROWEN
                                                                            Homo sapiens
                                                                                            05-MAR-1999;
                                                                                                  19-SEP-1994;
19-SEP-1995;
                            25-MAR-2004
                                                                                       17-0CT-2002
                                                                                                                                      Vbeta gene.
                      ADH70511;
           RESULT 991
ADH70511
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AAV2662-V26667 are primers used in the construction of plasmid pICNY which is used in a novel method of detecting the presence of a target PS112 polymucleotide in a test sample. The method can also be used to detect mRNA of PS112 in a test sample. The method can be used for diagnosis of prostate cancer, as the presence of PS112 is an indicator of prostate cancer. Antibodies against the polypeptides may be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate; disease; PS112 gene; detection; diagnosis; cancer; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting a target PS112 polynucleotide - used for diagnosing prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16; DB 1; Length 18; Pred. No. 1.1e+03;
                                              Score 16; DB 1; Length 17;
Pred. No. 1e+03;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 1 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
Sequence 17 BP; 0 A; 0 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gordon J, Hodges SC,
-Rapp L, Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. No. 1.1 tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 87; 104pp; English.
                                0.4%; Scc.
100.0%; Pre
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PS112 gene primer PS112.Fl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cohen M, Friedman PN, Gordon J
Kratochvil JD, Roberts-Rapp L,
                                                                                                                                                          2318 TGTGTGTGTGTGTG 2333
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                                                                                                                                                                                                                                                                                                                                           AAV26662 standard; DNA; 18 BP
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1D AAX77494 standard; DNA; 18 |

XX

XX AAX77494;

XX OS-AUG-1999 (first entry)
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Les 16; Conservative
                                                      Query Match 0.4
Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibody; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-240838/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ABBO ) ABBOTT LAB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-SEP-1998
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                                                                                                                                                                                                                                                                                                                                                                                                 AAV26662;
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                                                                                                                                                                                                                                                                                                RESULT 992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers associated diseases which comprises a panel of nucleic acid primers becilitacally priming and allowing amplification of each Vbetage gene, VbetaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases including autoimmune diseases, degenerative nervous system diseases, graft versus host disease, hypersenativity diseases, infectious disease, and neoplastic disease. Autoimmune diseases include Addison's disease, atrophic gastritis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivities such as those present in allorgies, Type II hypersensitivities such as those present in allorgies, Type II hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections manifested in leprosy. Infectious diseases include viral infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                            degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; hypersensitivity disease; arrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; HIV; fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                            T-cell associated disease; Wheta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 705; 164pp; English.
                                                                                                                                                                                                                                                                                                                           Human Vbeta gene repeat sequence #301.
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                                rererererere 16
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95US-00531241
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Klass MR; Stroupe SD;

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Gaps

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0; Indels

Synthetic

misc_RNA

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This invention describes a novel method for quantitating genetic instability independent of microsatellite alterations by treating a compatison pair comprising genomic DNA from tunour cells and genomic DNA from tunour cells and genomic DNA from normal cells. The method involves the cells from the same individual with oligonucleotide primers selected from adamine and same individual with oligonucleotide purimes selected from adamine and x = 3.7 (3) a nucleotide sequence (CG)xRY, where R is as in (1) and X is a pyrimidine selected from cytosine, thymine, and uracil and x = 3.7, (iv) a nucleotide sequence (CG)xRY, where R is as in (1) and x = 3.7, (iv) a nucleotide sequence (CG)xRY, where R is a pyrimidine selected from cytosine, thymine, and uracil and x = 3.7, (v) a nucleotide sequence (CA)xRY, where R is a purime selected from cytosine, (vi) a nucleotide sequence (CA)xRY, where R is a purime selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where R is a purime selected from adenine and guanine and Y is a pyrimidine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where R is a purime selected from adenine and continue selected from adenine and via a purime selected from adenine and via a purime selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, and (ix) a combination of the primers. The method is useful for detecting genomic instability which are commonly associated with the various stages of neoplastic transformation and carcinogenesis. The method is rapid and simple
                    Primer, quantitation; genetic instability; tumour cell; detection; neoplastic transformation; carcinogenesis; DNA/RNA hybrid; ss.
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                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Col 31-32; 27pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Quantitating genetic instability.
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"uracil"
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Best Local Similarity 100.0
Matches 16; Conservative
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/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HEAL-) HEALTH RES INC
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                                                                                                            Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Instanting and comparison of microsareline alterations by treating a comparison pair comporising genomic DNA from tumour cells and genomic DNA from normal cells. The method involves the cells from the same individual with Oligonucleotide sequence (CG) xRG, where R is a purine selected from adenine and guanine and x = 3-7, (ii) a mucleotide sequence (CG) xRY, where R is as in (i) and Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (iii) a nucleotide sequence (CG) xRY, where R is as in (i) and X = 3-7, (iv) a nucleotide sequence (CG) xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (v) a nucleotide sequence (CG) xRY, where R is a purine selected from cytosine, thymine, and uracil, and x = 6-16, (vi) a nucleotide sequence (CA) xRY, where R is a purine and guanine and X = 6-16, (vii) a nucleotide sequence (CA) xRY, where R is a purine selected from adenine and guanine and x = 6-16, (vii) a nucleotide sequence (CA) xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA) xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA) xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA) xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, and (ix) a combination of the primers. The method is useful for detecting genomic instability which are commonly associated with the various stages of neoplastic
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                                                                              Primer, quantitation; genetic instability; tumour cell; detection; neoplastic transformation; carcinogenesis; DNA/RNA hybrid; ss.
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100.0%; Pred. No. 1.1e+03;
:ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                     Location/Qualifiers
17. .18
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Les 16; Conservative
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                          USS912147 primer 38.
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Gaps

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16

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Query Match

Best Loca Matches

AAX77493;

AAX77493/ ID AAX7 XX AAX7 XX DT 05-A XX XX

/note= "uracil"

neoplastic transformation; carcinogenesis; ss

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This invention describes a novel method for quantitating genetic instability independent of microsatellite alterations by treating a comparison pair comprising genomic DNA from tumour cells and genomic DNA from normal cells. The method involves the cells from the same individual cutton normal cells. The method involves the cells from the same individual control of primers selected from adenine and selected from adenine and quanine and x = 3-7 (ii) a nucleotide sequence (CG)xRX, where R is as in (i) and Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (iv) a nucleotide sequence (CG)xXY, where R is as in (i) and X = 3-7, (iv) a nucleotide sequence (CG)xYY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (v) a nucleotide sequence (CG)xXY, where R is a pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (vi) a nucleotide sequence (CA)xRY, where R is a purine selected from adenine and guanine and Y is a pyrimidine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where R is a pyrimidine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, and (ix) a combination of the primers The method is useful for detecting genomic instability which are commonly associated with the various stages of neoplastic transformation and carcinogenesis. The method is rapid and simple
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0.4%; Score 16; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1.1e+03;
Matches 16; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Quantitating genetic instability.
                                                                                                                                                                                                                                                                                                                          Stoler D, Basik M;
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/note= "uracil"
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AAX77492 standard; DNA; 18 BP.
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                                                     USS912147-A.
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100.0%; Pred. No. 1.1e+03;
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RESULT 996

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This invention describes a novel method for quantitating genetic instability independent of microsatellite alterations by treating a comparison pair comprising genomic DNA from thmour cells and genomic DNA from thmour cells and genomic DNA from thmour cells and genomic DNA from thmour cells. The method involves the cells from the same individual with oligonucleotide primers selected from defining and guanine and x = 3-7 (dix) a nucleotide sequence (CG)xRX, where R is as in (i) and X is a pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (iii) a nucleotide sequence (CG)xRX, where R is as in (i) and X is a nucleotide sequence (CG)xRX, where R is a purine selected from cytosine, thymine, and uracil and x = 3-7, (v) a nucleotide sequence (CG)xRX, where R is a purine selected from definine and guanine and X is a pyrimidine selected from definine and guanine and Y is a pyrimidine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRX, where R is a purine selected from adenine and guanine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRX, where R is a pyrimidine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRX, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, and (ix) a combination of the primers. The method is useful to detecting genomic instability which are commonly associated with the various stages of neoplastic transformation and carcinogenesis. The method is rapid and simple
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neoplastic transformation; carcinogenesis; ss.
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                                                                                                                                                                               Quantitating genetic instability.
                                                                           Basik M;
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                                                                           Anderson G,
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100.0%; Pred. No. 1.1e+03;
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tive 0; Mismatches
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                                  Claim 4; Col 29-30; 27pp; English
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Quantitating genetic instability
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Best Local Similarity 100.'
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-357197/30.
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trom normal cells. The method involves the cells from the same individual with oligonucleotide primers selected from deanine and sequence (CG)xRG, where R is a purine selected from adenine and sequence (CG)xRG, where R is as in (1) and Y is a complexity of a nucleotide sequence (CG)xRY, where R is as in (1) and x = 3-7, (1ii) and coloride sequence (CG)xRY, where R is as in (1) and x = 3-7, (1ii) and coloride sequence (CG)xRY, where R is as in (1) and x = 3-7, (1iv) a conclectide sequence (CG)xRY, where R is a purine selected from cytosine, thymine, and uracil and x = 3-7, (v) a nucleotide sequence (CG)xRY, where R is a purine selected from deanine and guanine and Y is a pyrimidine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where R is a purine selected from adenine and guanine selected from denine and guanine selected from adenine and guanine selected from adenine and guanine selected from cytosine, thymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, (vii) a nucleotide sequence (CA)xRY, where Y is a pyrimidine selected from cytosine, thymine, and uracil and x = 6-16, and (ix) a combination of the primers The method is useful for detecting genomic instability which are commonly associated with the various stages of neoplastic transformation and carcinogenesis. The method is rapid and simple
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 16; DB 1; Length 18; [00.0%; Pred. No. 1.1e+03; [ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 9 A; 8 C; 0 G; 1 T; 0 U; 0 Other;
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/note= "uracil"
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nes 16; Conservative
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Matches
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(CG)xRG, where R is a purine selected from adenine and guanine and x = 3-

(i) a nucleotide sequence (CG)xRY, where R is as in (i) and Y is a

pyrimidine selected from cytosine, thymine, and uracil and x = 3-7, (iii)

a nucleotide sequence (CG)xRR, where R is as in (i) and x = 3-7, (iv) a

nucleotide sequence (CG)xRY, where Y is a pyrimidine selected from

Cytosine, thymine, and uracil and x = 3-7, (v) a nucleotide sequence

(CA)xRG, where R is a purine selected from adenine and guanine and x = 6-

16, (vi) a nucleotide sequence (CA)xRY, where R is a purine selected from

chymine, and uracil, and x = 6-16, (vii) a nucleotide sequence (CA)xRR,

where R is a purine selected from adenine and guanine and x = 6-16,

(vii) a nucleotide sequence (CA)xYY, where Y is a pyrimidine selected

CYCOSINE, thymine, and uracil and x = 6-16, and (ix) a combination

of the primers. The method is useful for detecting genomic instability

which are commonly associated with the various stages of neoplastic

transformation and carcinogenesis. The method is rapid and simple
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic, gene therapy, PS112; recombinant expression system, PS112 epitope, prostate disease, tumours, metastasis, predisposition,
                                                                                                                                                                                                                                                                                                                                                          Gaps
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Stroupe SD,
                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 16; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 1.1e+03; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                       Sequence 18 BP; 9 A; 8 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nan PN, Gordon J, Hodges SC,
Roberts-Rapp L, Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PS112 sequencing primer seqid 15.
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                                                                                                                                                                                                                                                                                                                                                                                         2318 TGTGTGTGTGTGTG 2333
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97US-00946869.
99US-00418887.
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KRATOCHVIL J D.
ROBERTS-RAPP L.
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FRIEDMAN P N.
GORDON J.
HODGES S C.
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STROUPE S D.
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08-OCT-1997;
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(RUSS/)
(STRO/)
(YUHH/)
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(KLAS/)
(KRAT/)
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(GORD/)
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The invention describes a method of detecting the presence of a target PS112 polynucleotide in a test sample. The method comprises: contacting the test sample with at least one PS112-specific polynucleotide or its complement; and detecting the presence of the target PS112 polynucleotide in the test sample, where the PS112-specific polynucleotide has at least complements or complements. Also described are: detecting mRNA of PS112 in test sample; a test kit, useful for detecting PS112 polynucleotide in test sample; a test kit, useful for detecting PS112 polynucleotide in test sample; a test kit, useful for detecting PS112 polynucleotide in test sample; a test kit, useful for detecting PS112 polynucleotide in test sample; a test kit, useful for a fragment derived from a PS112 control sequence compatible with a desired host, where the nucleic acid sequence of sequence has at least 50% identity to a sequence of SEQ ID NOS: 1-10, or their fragments or complements; a cell transfected with the recombinant expression system convibence acid sequence comparising at least 50% identity to a sequence comparises SEQ ID NOS: 1-10, or their fragments or complements; a cell transfected with the recombinant expression system complements; a cell sample or its fragment or their fragments of a PS112 polynucleotide or its fragments or or has at least 50% identity with SEQ ID NOS: 2-10, or their complements, or has at least 50% identity with SEQ ID NOS: 9 or 10. The method is DNA having at least 50% identity with SEQ ID NOS: 9 or 10. The method is complements, or has at least 50% identity with SEQ ID NOS: 9 or 10. The method is complement for detecting the presence of a target PS112 polynucleotide in a test sample. The methods, test kit, polynucleotides and polypeptides, and a propositicating, preventing and traating prostate disasses, tumours or metal for detecting, preventing and traating prostate disasses, tumours or metal for detecting the presence of a target PS12 polynucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prognosticating, preventing and treating prostate diseases, tumours metastases or in determining the predisposition of an individual to diseases and conditions of the prostate. e.g. prostate cancer. This sequence represents a primer used to sequence PS112 polynucleotides.
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1.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 1 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
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100.0%; Pred. No. 1...
0; Mismatches
                       2; SEQ ID NO 15; 53pp; English.
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Matches 16; Conservative
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                         Example
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13-FEB-2003.

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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous cassociated with development or maintenance of chronic myelogenous leukemia, (CML), promyelocytic leukemia, Burkitt's lymphoma, or acute leukemia, CML), promyelocytic leukemia, breaat cancer, colon carcinoma, neuroblastoma and lung cancer. Cleukemia, breaat cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant care directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a confine the may also be formed in the motif of a hairpin, chapatitis delta virus, group I intron or RNaseP-like RNA. These ribozymes can be used to inhibit the development or expression of a transformed corresponding spene. Cleavage of target mRNAs expression of the corresponding spene. Cleavage of target mRNAs expression of the motific and transformed cells elicits inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of drug theraples for tumours. The ribozymes may also be used to study cancer and mutations within cells. (Updated on 25-MAR-2003 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding multiple drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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0.4%; Score 16; DB 1; Length 19;
Best Local Similarity 75.0%; Pred. No. 1.2e+03;
Matches 12; Conservative 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 8 A; 1 C; 6 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Fig 10; 69pp; English.
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92US-00882885.
92US-00882885.
92US-00936410.
92US-00936421.
92US-00936531.
92US-00936531.
92US-00946532.
93US-00066123.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    delta chain PCR primer #1
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AGAUGAUGAAGAUGAU 17
                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                             Draper KG;
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1993-386203/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           correct PN field.)
                                                                                                                                                                                                                                                                                                                                                               rhompson JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
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                            14-MAY-1992;
26-AUG-1992;
                                                                                  26-AUG-1992;
26-AUG-1992;
26-AUG-1992;
                                                                                                                                                                                                                       19-JAN-1993;
                                                                                                                                                                 26-AUG-1992
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              resistance.
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WO2003011337-A1

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                                                                                                                                                                                                                                                                                                     The invention relates to a drug composition containing as an active component a substance which activates Fc receptor gamma chain. Also included are detecting oligodendroglia and their precursor cells capable of forming myelin (using as an indicator the expression of Fc receptor gamma chain in the precursor cells), investigation of the expression of Fc receptor gamma chain in animal brain tissue (by immune typing, cytochemical analysis, gene amplification or Western blotting) and kits for these methods, containing anti-Fc receptor gamma chain antibody or amplification primers. The drug is used for treatment and prevention of neurodegenerative diseases, and disorders of myelin formation, such as multiple sclerosis, Krabbe's disease, adrenoleukodystrophy and metachromic leukodystrophy. The present sequence is a CD3 delta chain PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              non-human animal model; demyelinating disease; myelinogenesis inhibition; myelinogenesis signal molecules; oligodendroglia; screening; myelin growth regulator; multiple sclerosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Non-human animal model for demyelinating disease-related PCR primer #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                         Drug compositions containing Fc receptor gamma chain activator for treatment and prevention of neurodegenerative disorders including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel non-human animal model for demyelinating disease in which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   metachromic leukodystrophy. The present sequence is primer used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 16; DB 1; Le
100.0%; Pred. No. 1.2e+03;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                 Example 9; Page 49; 109pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; PER
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                                                                                                                                                Aiso S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13
                                             22-JUL-2002; 2002WO-JP007378
                                                                            30-JUL-2001, 2001JP-00229553
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                Asou H,
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                                                                                                                                                                                 WPI; 2003-248118/24.
                                                                                                                                                                                                                                                 multiple sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                               (UYKE-) UNIV KEIO
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                                                                                                                                                Nakahara J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC49455;
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PSII2 polynucleotide in a test sample. The method can also be used to detect mRNA of PSII2 in a test sample. The method can be used for diagnosis of prostate cancer, as the presence of PSII2 is an indicator of prostate cancer. Antibodies against the polypeptides may be used as markers, or to treat prostate cancer
                                                                                                 The invention comprises a non-human animal model for demyelinating disease - in which myelinogenesis is inhibited by a defect of myelinogenesis signal molecules in oligodendroglia. The non-human animal model of the invention is useful for screening for a myelin growth regulator, or for screening for a therapeutic agent which is useful for treating a demyelinating disease such as multiple sclerosis. The present DNA sequence represents a PCR primer that was used in an example of the
myelinogenesis is inhibited by defect of myelinogenesis signal molecules in oligodendroglia, for screening for therapeutic agent for multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate; disease; PS112 gene; detection; diagnosis; cancer; treatment; antibody; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting a target PS112 polynucleotide - used for diagnosing prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV26652-V26667 are primers used in the construction of plasmid pICNY which is used in a novel method of detecting the presence of a target
                                                                                                                                                                                                                                                                                     Ouery Match 0.4%; Score 16; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 1.2e+03; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gordon J, Hodges SC,
-Rapp L, Russell JC,
                                                                   Example; SEQ ID NO 3; 56pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 88; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PS112 gene primer PS112.R3
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                                                                                                                                                                                                                                                                                                                                                            2303 CACAGAGCTTTGGTCT 2318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ABBO ) ABBOTT LAB.
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                                 sclerosis.
                                                                                                                                                                                                                          invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV26667;
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Klass MR; Stroupe SD;

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The invention relates to antisense oligonucleotides targeted to a nucleic acid molecule encoding human Survivin, where the antisense oligonucleotide inhibits the expression of human Survivin. These antisense oligonucleotides are used in the treatment of an animal suffering from a disease or condition associated with Survivin, e.g. a hyperproliferative condition such as cancer, and comprises administering a therapeutically or prophylactically effective amount of the antisense oligonucleotides can also be used to treat a human suffering from a disease or condition characterised by a reduction in apoptosis comprising administering the antisense oligonucleotide to a human. In addition, the antisense oligonucleotide to a human. In addition, the call cycle, or inhibit the proliferation in a cancer cell by contacting the cell vith the antisense oligonucleotide. AAS215A1-AAS21768 represent Survivin nucleic acids, and antisense oligonucleotides targeted to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel antisense compounds for modulating the expression of Survivin and treatment of cancer.
                              Gaps
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                                                                                                                                                                                                                                                                                                      Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis; 
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                            Indels
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           Pred. No. 1.2e+03;
Mismatches 0;
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                                                                                                                                                                                                                                                                        Mouse Survivin antisense oligonucleotide #57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 18; Page 62; 120pp; English.
100.0%; Pre-
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                                                            832 TGGCTGGTGCTGC 847
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                                                                                                                                                                         AAS21755 standard; DNA; 20
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hes 16; Conservative
                              Conservative
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             Local Similarity
nes 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                        Mus musculus.
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                                                                                                                                                                                                                                         21-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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Matches
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Gaps ö

0.4%; Score 16; DB 1; Length 20;

Query Match

ABT07496/c

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human neuroblastoma with good prognosis. The DNA sequences of the invention are useful in clarifying good/poor prognosis of neuroblastoma. The present DNA sequence was used in the exemplification of the invention
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100.0%; Pred. No. 1.2e+03;
ive 0; Mismatches 0; Indels
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                                                                                                                                                         Neuroblastoma; prognosis; ds; oligonucleotide.
                                                                          Neuroblastoma-related DNA sequence #71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 23(1); 444pp; Japanese.
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24-AUG-2001; 2001JP-00255226.
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24-AUG-2001; 2001JP-00255260.
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22-SEP-2003 (first entry)
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nes 16; Conservative
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                                                                                                                                                                                                                                        Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel compound 8-50 nucleotides in length targeted to a nucleic acid molecule encoding a protein phosphatase 2 catalytic beta subunit, where the compound specifically hybridises with and inhibite the expression of protein phosphatase 2 catalytic beta subunits, or specifically hybridises with at least an 8-nucleotide compounds and active site on a nucleic acid molecule encoding a protein phosphatase 2 catalytic beta subunit and for treating diseases or conditions associated with computating the expression of protein phosphatase 2 catalytic beta subunits and for treating diseases or conditions associated with a subunits and for treating diseases or conditions associated with a berrant insulin regulation or diabetes or a hyperproliferative disorder, particularly cancer. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and in the distinguishing between functions of various members of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                 protein phosphatase 2 catalytic beta subunit; antisense compound; cancer; hyperproliferative disorder; diabetes; inflammation; tumour; rat; ds.
                                                                                                                                                                                                                                                                                                                                                       Cytostatic; antidiabetic; antisense therapy; aberrant insulin regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides targeted to nucleic acid encoding Protein Phosphatase 2 catalytic subunit beta, useful for treating diseases related to Protein Phosphatase 2 catalytic subunit beta expression, such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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100.0%; Pred. No. 1.2e+03;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                             Rat protein phosphatase 2 oligo inhibitor SEQ ID No 110.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 16; Page 98; 137pp; English.
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                             ABT07496 standard; DNA; 20 BP
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                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-657588/70.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200264737-A2
                                                                                                                                                                                              14-NOV-2002
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                                                                                                               ABT07496;
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Gaps

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Neuroblastoma; prognosis; spontaneous regression; primer; probe; ds;
                                  Neuroblastoma~related oligonucleotide #88.
                                                                                                                              (CHIB-) CHIBA PREFECTURE.
(HISM ) HISAMITSU PHARM CO LTD.
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ABT43156

ABT43156 ID ABT4 XX AC ABT4 XX

RESULT 1009

Matches

ઠ g Length 20;

Query Match

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human neuroblastomas with poor prognosis over those with a good prognosis. The nucleic acids of the invention are useful as a tool for distinguishing neuroblastomas with a favourable prognosis (spontaneous regression) from neuroblastomas with a poor prognosis (high malignancy). The DNA sequences ABT32224 - ABT32571 represent oligonucleotides used in an example of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence repeat; ISSR; SSR; PCR; primer; genotyping; plant;
                                              Nucleic acids having higher expression in human neuroblastoma with poor prognosis for diagnostic prediction of neuroblastoma prognosis.
                                                                                                         The invention comprises nucleic acids that show increased expression in
                                                                                                                                                                                                                             0.4%; Score 16; DB 1; Length 20;
100.0%; Pred. No. 1.2e+03;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                      Sequence 20 BP; 4 A; 8 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA FINGERPRINTING & DIAGNOSTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                        anchored (ISSR)-PCR primer - SEQ ID 26.
                                                                                 Example 5; Page 26; 111pp; Japanese.
                                                                                                                                                                                                                                                                             2632 CCACATGTCCAGCACC 2647
                                                                                                                                                                                                                                                                                                                                                                ADD69468 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                            CCACATGTCCAGCACC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                     16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              animal; Basmati rice;
                                                                                                                                                                                                                                Query Match
Best Local Similarity
                       WPI; 2003-140476/13.
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 Nakagawara A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nagaraju JG;
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New set of inter-simple sequence repeats (ISSR)-PCR primers for genotyping eukaryotes, useful for genotyping diverse genomes of plant and

Claim 1; SEQ ID NO 26; 60pp; English.

genotyping euka animal systems.

The invention relates to a novel set of inter-simple sequence repeats (ISSR)-PCR primers for genotyping eukaryotes. The primers of the invention may be useful for genotyping diverse genomes of plant and animal systems, in particular for distinguishing Basmati rice varieties from non-Basmati rice varieties and traditional Basmati rice varieties from evolved Basmati rice varieties. The current sequence is that of the 3' anchored (ISSR)-PCR primer of the invention.

Sequence 20 BP; 1 A; 2 C; 8 G; 9 T; 0 U; 0 Other;

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, 5' or 3' end genomic flanking regions, 5' or and 1' intron-exon junctions or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or casal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinon: A composition of the invention has antiinflammatory, antiallergic, antiaethmatic, hypotensive, cuse in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense year therapeutic respiratory effect of an extiniflammatory steroid in a subject, for reducing or depleting levels of for entainflammatory seroid in a subject, for reducing levels of adenosine cofficion producing bronchodilation, increasing levels of adenosine lung suffactant in a subject stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiinsthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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                                      Indels
ore 16; DB 1; Le
red. No. 1.2e+03;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 7398; 872pp; English
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S;
  0.4%; Score 16;
100.0%; Pred. No.
  100.0%; Pr.
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L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                     Human oligonucleotide sequence.
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                                        Conservative
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                    Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200285308-A2.
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R78585-derived oligonucleotide SEQ ID 7398.
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Tang L, Shahabuddin
                                                                                                             2980 ACCAGGCTTTTTCTG 2995
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                                                     16; Conservative
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                       Best Local Similarity
Matches 16; Conserv
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Miller S,
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Query Match
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inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary dysperension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chimeric; antisense oligonuclectide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPcES-1; mPcES-1; inhibitor; microsomal prostaglandin E2 synthase; inhibitor; crytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; immunomodulatory; cardiant; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1374.
                                                                                                                                                                                                                                                                                                      0.4%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 1.2e+03; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                          Seguence 20 BP; 8 A; 4 C; 5 G; 3 T; 0 U; 0 Other;
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'note= "2'-0-methocyethyls"
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ADM15187 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery chartuctions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction,
                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity, lung, cancer, surfactant depletion, antiallergic, antiinflammatory, antiasthmatic,
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nucleic acids associated with lung airway or lung dysfunction, and
bronchodilating agent.
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  0.4%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 1.2e+03; ive 0; Mismatches 0; Indels
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The present sequence represents a chimeric antisense oligonucleotide charged to human microsomal prostaglandin E2 synthase (mpcBS-1). The human mPGES-1 gene is located on chromosome 9, more specifically to operate on the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal carrisense oligonucleotides and antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamaticy, neuroprotective, nootropic, antidiamaticy, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound to be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; s.
New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                  Claim 4; SEQ ID NO 1374; 132pp; English
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Matches 16; Conservative
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ADM15240/c
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPCES-1). The human mPCES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPCES-1, which specifically whybridise with the nucleic acid encoding inhibite its expression; (2) a method of inhibiting the expression of mPCES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPCES-1. MPCES-1 chimeric antidabetic, immunomodulator, cardiant, neuroprotective, cantidiabetic, immunomodulator, and cardiovascular activities, and can antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mPCES-1 inhibitors and in gene therapy. The antisense compound to used as mPCES-1 inhibitors and in gene therapy. The antisense compound condition associated with mPCES-1 e.g., inflammation, Alzheimer's condition associated with mPCES-1 e.g., inflammation, Alzheimer's chiscase, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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PS112 epitope; prostate disease; tumours; metastasis; predisposition;
prostate cancer; PCR; primer; ss.
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/note= "2'-0-methoxyethyls"
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                                                                                                                                                        25-SEP-2002; 2002US-0413549P.
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ADQ96632 standard; DNA; 20
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                                     WO2004028458-A2
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(KLAS/) (KRAT/) (FRIE/) (GORD/) (HODG/) COHE/)

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This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active on a gene vitel for cell growth or viablity, and where the gene is subject to loss of heterozygosity (LCH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene specific inhibitor (ASI) targeted to an allele of a first essential gene specific inhibitor of the precancerous condition, where the normal somatic calls of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene comment in a population and targets only one allelic forms present in the cused in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic cancers, benigh tumours, endometriosis, polycystic kindry disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                   Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Housman D, Ledley FD, Stanton VP;
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                                                                                                                            Human polymorphic region 884.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 GTGAAGATGCTGAAAG
                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (VARI-) VARIAGENICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-521232/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                           WO9841648-A2
                                                                          30-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG14272;
                           AAZ26695;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG14272
SXXXE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ઢ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a method of detecting the presence of a target PS112 polynucleotide in a test sample. The method comprises: contacting the ps112 polynucleotide or lies complement; and detecting the presence of the target PS112 polynucleotide or lies complement; and detecting the ps112-specific polynucleotide or lies in the test sample, where the PS112-specific polynucleotide has at least in the test sample, where the PS112-specific polynucleotide has at least complements or complements. Also described are: detecting mRNA of PS112 in treagments or complements. Also described are: detecting mRNA of PS112 in test sample; a purified polynucleotide or fragment derived from a PS112 cent a recombinant expression system comprising a nucleic acid sequence that includes an open reading frame derived from PS112 oplynucleotide or fragment derived from a PS112 central sequence compatible with a desired from PS112 oplynucleotide or complements; a cell transfected with the recombinant expression system or with a nucleic acid sequence encoding at least one pS112 epitope, where the nucleic acid sequence encoding at least one pS112 epitope, where the nucleic acid sequence comprises SEQ ID NOS: 1-10, or their fragments or complements, a central polynucleotide or its fragment, where the polynucleotide has at least 50% identity with SEQ ID NOS: 2-10, or their comprising a plant fragments or complements or complements or detecting the presence of starget PS112 polynucleotide in a test sample. The methods, test kit, polynucleotides and polypeptides, and test sample. The methods, test kit, polynucleotides and polypeptides, and tests as an individual to metastases or in determining the predisposition of an individual to metastases or in determining the predisposition of an individual to cequence represents a primer used to sequence eg. prostate cancer. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting a target PS112 polynuclectide, useful in diagnosing, staging, monitoring, prognosticating, preventing and treating prostate cancer, comprises contacting the test sample with PS112-specific polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ϋ́υ
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Klass MR;
Stroupe SD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.4%; Score 16; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.2e+03;
Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cohen M, Friedman PN, Gordon J, Hodges SC,
Kratochvil JD, Roberts-Rapp L, Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; SEQ ID NO 20; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        832 TGGCTGGTGGTGC 847
                                                          96US-00727688.
97US-00946869.
99US-00418887.
     22-JAN-2004; 2004US-00763992
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                                                                                                                                                                                                                                                                      KLASS M R.
KRATOCHVIL J D.
ROBERTS-RAPP L.
                                                                                                                                                                   COHEN M.
FRIEDMAN P N.
                                                                                                                                                                                                                    GORDON J.
HODGES S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-479676/45.
                                                                                                                                                                                                                                                                                                                                                   RUSSELL J C.
STROUPE S D.
                                                          08-OCT-1996;
08-OCT-1997;
15-OCT-1999;
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Gaps

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DB 1; Length 21; 1.3e+03; 0; Indels

26-FEB-2004 (first entry)

AAZ26695 standard; DNA; 21 BP.

RESULT 1017 AAZ26695 ID AAZ2

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Chou Q,

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The sequences given in AAQ27805-7 are primers which are complementary to intronic sequences within the amyloid precusor protein gene (APP). They were used to amplify exon 17. APP encodes various isoforms which are precursors of beta-amyloid protein. The beta-amyloid protein is an approx. 4kD protein (39-42 amino acids) which is an internal cleavage fooduct from APP. There are five distinct isoforms of APP containing 563, 695, 714, 751 and 770 amino acids. These are generated by alternative splicing of primary transcripts of APP which is located on human (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
                                                                                                                                   Polynucleotide probe comprising nucleic acid encoding codon 717 mutant of human APP770, useful for determining genetic pre-disposition to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kanagawa T, Kamagata Y, Kurata S, Yamada K, Yokomaku T;
Furusho K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Concentration, quantification, mutation detection, polymorphic, polymerase chain reaction, PCR; ss.
                                                        Owen MJ, Mullan MJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 8 A; 3 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 16; DB 1; Le
100.0%; Pred. No. 1.4e+03;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic deoxyribooligonucleotide poly d.
                                                        Goate AM,
                (UNLO ) IMPERIAL COLLEGE SCI TECHN MED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NIBI-) JAPAN BIOINDUSTRY ASSOC.
(AGEN ) AGENCY OF IND SCI & TECHNOLOGY.
                                                                                                                                                                                                                  Disclosure; Page 31; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 21; 55pp; English
                                                        Chartier-Harlin MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3365 AAATTCTTCTAATTGC 3380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL56891 standard; DNA; 30 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99JP-00111601.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-657765/64.
                                                                                                                                                                              Alzheimer's disease.
                                                                                               WPI; 1992-284654/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-APR-1999;
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Koyama O,
                                                        Hardy JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL56891;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1020
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting the production of a primer extension product comprises using a fluorescence energy transfer (FET) labeled oligonucleotide that includes a 3'-5' exonuclease resistant quencher domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                quencher domain; subjecting the mixture to primer extension reaction conditions; and detecting a change in a fluorescent signal from the FET labelled probe. The method is useful for real-time monitoring of PCR amplification reactions. The present sequence was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a method (M1) for detecting the production of a primer extension product. M1 comprises: producing a primer extension mixture that includes a nucleic acid polymerase having 3'-5' exonuclease activity and a fluorescence energy transfer (FET) labelled oligonucleotide that includes a 3'-5' exonuclease resistant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beta-amyloid protein; amyloid precursor protein; isoform; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 16; DB 1; Length 21;
100.0%; Pred. No. 1.3e+03;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 3 A; 3 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 1; 52pp; English
                Human bcl-2 gene PCR primer, BCL-F1.
                                                                                                                                                                                                                                                                                                                                                             Spasic D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1873 GTGGAGGAGCTCTTCA 1888
                                                        Human; bcl-2; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВР
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                                                                                                                                                                                                                    25-FEB-2003; 2003WO-US005641
                                                                                                                                                                                                                                                           27-FEB-2002; 2002US-00087229
15-AUG-2002; 2002US-00222943
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                                                                                                                                                                                                                                                                                                                     (BIOS-) BIOSOURCE INT INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.43
Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   method of the invention.
                                                                                                                                                                                                                                                                                                                                                         Cabradilla CD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ27807 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APP exon 17 primer 5.
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-803791/75.
                                                                                                                                       WO2003072051-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9213069-A1
                                                                                                 Homo sapiens
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28-AUG-1991;
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28-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JAN-1992;
                                                                                                                                                                            04-SEP-2003.
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AAQ27807;

RESULT 1019

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The invention relates to the determination of the concentration of a nucleic acid target, using a fluorescently labeled probe which produces reduced fluorescence emission when hybridised to the target nucleic acid. The method comprises meaturing the reduction in emission caused by the method comprises meaturing the reduction in emission caused by a real-time polymerase chain reaction, e.g. for nucleic acids by a real-time polymerase chain reaction, e.g. for quantifying microbial cells in co-cultures or symbiotic systems, for detecting gene mutations or polymorphisms, and for analysing melting curves of target nucleic acids to determine a Tm value. Methods of the invention allow target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and on materials are introduced that inhibit amplification by Taq polymerase (so conventional PCR conditions can be used). The specificity of PCR is kept high (amplification of primer dimers is delayed), and the limit of quantitation is reduced. Complex probes are not needed, and amplification can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation coefficient than conventional graphs so more accurate quantitation is possible. The current sequence represents a synthetic quantitation is decaying the base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kanagawa T, Kamagata Y, Kurata S, Yamada K, Yokomaku T; Furusho \mathbb{K}_i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Concentration, quantification, mutation detection, polymorphic, polymerase chain reaction, PCR, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 16; DB 1; Length 30; 9.2%; Pred. No. 1.8e+03; ve 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic deoxyribooligonucleotide poly b.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NIBI-) JAPAN BIOINDUSTRY ASSOC.
(AGEN ) AGENCY OF IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3474 ATATATATTATTATTGAGTTTTT 3497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               selectivity of a target nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABL56889 standard; DNA; 30 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-APR-2000; 2000EP-00108643.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.4%;
Best Local Similarity 79.2%;
Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-JUL-2002
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Koyama O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.

WPI; 2000-657765/64.

The invention relates to the determination of the concentration of

Example 5; Page 21; 55pp; English

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Gaps

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The method comprises measuring the reduction in emission caused by hybridisation. The new method is particularly used to quantify target nucleic acids by a real-time polymerase chain reaction, e.g. for nucleic acids by a real-time polymerase chain reaction, e.g. for quantifying microbial cells in co-cultures or symbiotic systems, for detecting gene mutations or polymorphisms, and for analysing melting curves of target nucleic acids to determine a Tm value. Methods of the invention allow target nucleic acids to be quantified quickly, easily and conventional per nucleic acids to be quantified quickly, easily and materials are introduced that inhibit amplification by Tag polymerase (so conventional PCR conditions can be used). The specificity of PCR is kept high (amplification of primer dimers is delayed), and the limit of quantitation is reduced. Complex probes are not needed, and amplification can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation coefficient than conventional graphs so more accurate quantitation is possible. The current sequence represents a synthetic decorate decorate passe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Measurement of nucleic acids, using a nucleic acid probe and analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorecence of the fluorochrome when hybridised with a rarget nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
nucleic acid target, using a fluorescently labeled probe which produces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     % Match 0.4%; Score 16; DB 1; Length 30; Local Similarity 79.2%; Pred. No. 1.8e+03; les 19; Conservative 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; fluorochrome; nucleic acid probe; fluorescence.
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'KANK-) KANKYO ENG KK.
'KEIZ-) KEIZAI SANGYOSHO SANGYO GIJUTSU SOGO KEN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3497
                                                                                                                                                                                                                                                                                                                                                                                               selectivity of a target nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Page 17; 34pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3474 ATATATATATATTTGAGTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 ATATATTTTTTTTTTTTTTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BIOI-) BIOINDUSTRY KYOKAI SH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-APR-1999; 99JP-00111601.
24-AMG-1999; 99JP-0023666.
30-AMG-1999; 99JP-00242693.
01-FBB-2000; 2000JP-00028996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-APR-2000; 2000JP-00120097.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Poly b nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA97613 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-134193/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2001286300-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA97613;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1022
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Matches
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Yamada K;

Kurata S,

Torimura M,

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The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleotide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                 Fluorescently-labeled nucleic acid probes for assaying nucleic acids and their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and mirobbial identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Probe; polymorphism detection; mutation detection; disease diagnosis; microbial identification; ss.
                            Probe; polymorphism detection; mutation detection; disease diagnosis; microbial identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match

0.4%; Score 16; DB 1; Length 30;
Best Local Similarity 79.2%; Pred. No. 1.8e+03;
Matches 19; Conservative 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                           (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Probe poly d for assaying nucleic acids.
Probe poly b for assaying nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3474 ATATATATATTTTATTGAGTTTTT 3497
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 12; Page 60; 152pp; Japanese.
                                                                                                                                                                                                                                                                                                         Kanagawa T, Kamagata Y,
                                                                                                                                                                  27-JUN-2001; 2001WO-IB001147.
                                                                                                                                                                                                27-JUN-2000; 2000JP-00193133.
                                                                                                                                                                                                              03-AUG-2000; 2000JP-00236115
26-SEP-2000; 2000JP-00292483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL95888 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-195876/25.
                                                                                                         WO200208414-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the invention
                                                                          Unidentified.
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                                                                                                                                       31-JAN-2002.
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                                                                                                                                                                                                                                                                                                                        Yokomaku T;
                                                                                                                                                                                                                                                                                                           Kurane R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL95888;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
                                                                            Gapa
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                                          Score 16; DB 1; Length 30;
Pred. No. 1.8e+03;
0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                        ss; fluorochrome; nucleic acid probe; fluorescence.
                 Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (KANK-) KANKYO ENG KK.
(KEIZ-) KEIZAI SANGYOSHO SANGYO GIJUTSU SOGO KEN
                                                                                                         3474 ATATATATATITATITGAGTTTTT 3497
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| ATATATATTTTTTTTTTT 24
                                                                                                                                    ATATATATTTTTTTTTTTTTT 24
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                                                                                                                                                                                                                ABA97615 standard; DNA; 30 BP
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24-AUG-1999; 99JP-00236666.
30-AUG-1999; 99JP-00242693.
01-FEB-2000; 2000JP-00028896.
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                                            Query Match
Best Local Similarity 79.2%;
Matches 19; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-APR-2000; 2000JP-00120097
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                                                                                                                                                                                                                                                                                                         Poly d nucleotide sequence
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                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                  JP2001286300-A
                                                                                                                                                                                                                                                                                                                                                                       Unidentified
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ABA97615
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Matches
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ABL95886
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AC ABL9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HPV infection; replication; cytostatic; virucide; cervical dysplasia; excinoma; oral mucosal cancer; laryngeal; vaccine; gene therapy; double internal trans-acting ribozyme; single; dITRz; CHOP portion; SNIPAA cassette; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3' flanking RNA of IT ribozyme (Rz) from CHOP portion of SNIPAA cassette.
                                                                                                                                                                                                                                 and
                                                                                                                                                 Yamada K;
                                                                                                                                                                                                                              Fluorescently-labeled nucleic acid probes for assaying nucleic acids their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 16; DB 1; Length 30; 79.2%; Pred. No. 1.8e+03; ive 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                               Kanagawa T, Kamagata Y, Torimura M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Christensen N, Thiboutot D;
                                                                                              (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY (KANK-) KANKYO ENG CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3474 ATATATAATTTATTGAGTTTTT 3497
                                                                                                                                                                                                                                                                                                              Example 12; Page 60; 152pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2001; 2001WO-IB001147
                             27-JUN-2000; 2000JP-00193133
03-AUG-2000; 2000JP-00236115
26-SEP-2000; 2000JP-00292483.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-2003; 2003WO-US020340.
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14-OCT-2002; 2002US-0417997P.
21-FEB-2003; 2003US-0449066P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI32816 standard; RNA; 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19; Conservative
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                                                                                                                                                                                               WPI; 2002-195876/25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
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                                                                                                                                                 Kurane R, I
Yokomaku T;
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ADI32816/c
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                                                                                                                                                                                  The invention relates to a novel method for treating Human papillomavirus (HPV) infection comprising administering to a patient a nucleic acid molecule that inhibits expression associated with HPV replication. The method of the invention has cytostatic and virucide applications and may be useful for treating HPV infections or HPV-associated conditions including cervical dysplasia, cervical carcinoma, oral mucosal cancer and laryngeal cancer. Furthermore, the method may be utilised during vaccine production and gene therapy. The current sequence is that of the HPV-targeted ribozyme (Rz)-related RNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor
                                                Treating human papilloma virus (HPV) infection, e.g. cervical dysplasias or HPV-associated cervical carcinoma, comprises administering to a patient a nucleic acid molecule that inhibits expression associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; compound; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 16; DB 1; Length 39; 88.8%; Pred. No. 2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compound simple sequence repeat primer (GT)7.5(AT)2.
                                                                                                                                                                                                                                                                                                                                                                            Sequence 39 BP; 32 A; 2 C; 2 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                             68.8%; Pred. .v.
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                                                                                                                                                Example 2; SEQ ID NO 56; 65pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 68.8
hes 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-277795/28.
               WPI; 2004-082869/08.
                                                                                                             HPV replication.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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The present invention describes an isolated, pure population of mammalian neuroepithelial stem cells, which are capable of self-renewal in adherent neuroepithelial stem cells, which are capable of self-renewal in adherent cell-independent (AFCI) culture medium and differentiation to central nervous system (CNS) neuronal or glial cells and to neuronal crest stem cells. Also described is an isolated population of mammalian CNS glial-restricted precursor (GRP) cells which can self-renew in the AFCI culture medium and can differentiate to CNS glial cells but not to CNS neuronal cells. The stem cells can be used to generate a population of mammalian motor neurons by incubating the stem cells in a medium comprises laminin-coated plates and NBP medium lacking chick embryo extract. The stem cells on also produce neural crest stem cells by inducing the cells to differentiate in vitro. The inducing step comprises replating the cells on a laminin-coated substrate and preferably cinducing a dorsalizing chick embryo extract. Inducing can also comprise adding a dorsalizing agent to the cells, preferably a bone morphogenetic protein (BMP) such as
                                                                                                                                                                                                                                                                                                                                                ö
directed primer, comprising a sequence complementary to an adaptor segment. The present sequence is an example of a compound SSR primer. The method represents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and germplasm comparisons
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuroepithelial stem cell; lineage restricted intermediate precursor; oligodendrocyte; self-renewal; neuron; diferentiation; neural crest cell; fibroblast growth factor; FGFR; receptor; CNS; central nervous system; glal cell; PCR primer; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                             Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                         Sequence 19 BP; 2 A; 0 C; 7 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                         Local Similarity 89.
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BMP-2, -4 or -7. The stem cells can be used to produce cells of the peripheral nervous system, by inducing the stem cells to differentiate in vitro to neural crest stem cells, and inducing these cells to differentiate. AAX00029 to AAX00054 represent PCR primers which are used in an example from the present invention to amplify different FGF and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ü
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA83415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
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                                                                                                                                             Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
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Best Local Similarity 89.5%; Pred. No. 1.2e+03;
Matches 17; Conservative 0; Mismatches
                                                                                                                Sequence 19 BP; 3 A; 10 C; 5 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclin B1 ribozyme binding site #134.
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                                                                                                                                                                                                                 1210 GGGGAGGGCTGCTTCGGCC 1228
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Best Local Similarity 89.5%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                              AAA85805 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       restenosis treatment
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                                                                                FGFR genes
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RESULT 1030

viral wart;

vivlemore401-10.rng

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antisebornheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and vincide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermaticis, actinic keratosis, acquamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diseases such as solviant proliferative eye diseases such as diseases such as electinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing correcting such as keloid, adhesion and hypertrophic or hypertrophic burn security such as keloid, adhesion and hypertrophic or hypertrophic burn security such as elected and such as the such as elected and such as keloid, adhesion and hypertrophic or hypertrophic burn and correcting and preventing and preventing secur. AMHSCOST regiments sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  short interfering nucleic acid; siNA; downregulation; inhibition; St stearoyl-COA desaturase; RNA interference; anorectic; antidiabetic; antiarteriosclerotic; cytostatic; virucide; obesity; diabetes;
                     antisickling; ophthalmological; keratolytic; gene therapy; vira atopic dermatitits; actinic keratosis; squamous cell carcinoma; assal cell carcinoma; seborrheic wart; virreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               exemplification of the present invention
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Best Local Similarity 89.5'
Matches 17; Conservative
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                                                                                                                                                                                        WO200130362-A2.
                                                                                                                              Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                26-OCT-1999;
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                                                                                                                                                                                    Mouse; lineage restricted precursor cell; neuron-restricted precursor; NRP; glial-restricted precursor; GRP; mouse neural tube; transplantation; antibody; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides populations of lineage-restricted precursor cells from mouse neural tube and mouse embryonic stem cells. These populations are of neuron-restricted precursor cells (NRPs) and glial-restricted precursor cells (GRPs). These cell populations are useful in the development of new transplant techniques, for transplantation in diseases where neuronal or glial degeneration has occurred, in the identification of drugs which enhance the survival and proliferation of transplanted cells, to identify genes specific to selected stages of development, and in the generation of cell-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New pure populations of neuron- or glial-restricted precursor cells and neuroepithelial stem cells from mouse neural tubes or embryonic stem cells for developing new drugs or techniques that enhance survival of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                 Murine lineage-restricted precursor cell population PCR primer #18.
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                           AAC88240 standard; DNA; 19 BP.
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                                                                                                           02-MAR-2001
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                                                                  AAC88240;
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Gaps

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The present invention describes a short interfering nucleic acid (siNA) by RNA interference. Also described: (1) modulating expression of the SCD (stearoyl-CoA desaturase) gene by RNA interference. Also described: (1) modulating expression of SCD genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA. SCD inhibiting siNAs have anorectic, antidiabetic, antiarteriosclerctic, cytostatic and virucide activities. The siNAs can be used to modulate expression of SCD genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double-stranded short interfering nucleic acid;
short interfering nucleic acid; siNA; downregulation;
swacular endothelial growth factor receptor; VEGFR; antiangiogenic;
cytostatic; antidiabetic; ophthelmological; antiarthritic; antipsoriatic;
nephrotropic; gynaecological; angiogenesis-associated condition; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human VEGFR1 short interfering nucleic acid (siNA) SEQ ID NO:158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New short interfering nucleic acid, useful e.g. for treatment diagnosis of obesity or diabetes, downregulates expression of stearoyl-CoA desaturase gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 15.8; DB 1; Length 19; 39.5%; Pred. No. 1.2e+03; lve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
  genetic engineering; pharmacogenomic; gene mapping; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                     Thompson J;
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                                                                                                                                                                                              20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
                                                                                                                                                                                                                                                       29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
20-SEP-2002; 2002US-0412304P.
                                                                                                                                                           13-FEB-2003; 2003WO-US004317.
                                                                                                                                                                                                                                                                                                                                       15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 89.5%;
nes 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-721687/68.
                                                                             WO2003070885-A2
                                                                                                                    28-AUG-2003
                                        Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         short interfering nucleic acid; siNA; downregulation; inhibition; SCD; stearcyl-CoA desaturase; RNA interference; anorectic; antidiabetic; antiarteriosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New short interfering nucleic acid, useful e.g. for treatment and diagnosis of obesity or diabetes, downregulates expression of the stearoyl-CoA desaturase gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stearoyl-CoA desaturase siNA oligonucleotide SEQ ID NO:447.
atherosclerosis; cancer; viral infection; drug screening;
                      genetic engineering; pharmacogenomic; gene mapping; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 4 A; 5 C; 7 G; 0 T; 3 U; 0 Other;
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3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 157; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                       Mcswiggen J, Beigelman L, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             784 TACACCTGCCTGGCGGCA 802
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29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
20-SEP-2002; 2002US-0412304P.
15-JAN-2003; 2003US-0412304P.
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                                                                                                                                                                                                                  2002US-0358580P.
2002US-0363124P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           73.78;
                                                                                                                                                                           13-FEB-2003; 2003WO-US004317
                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC
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nes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-721687/68.
                                                                                                WO2003070885-A2.
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11-MAR-2002;
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                                                                                                                                       28-AUG-2003
                                                           Synthetic.
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Gaps

double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; avacular endothelial growth factor receptor; VERFR; antiangiogenic; cytostatic; antidabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; alabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma;

20-FEB-2003; 2003WO-US005022.

WO2003070910-A2.

28-AUG-2003

Homo sapiens.

Synthetic

2002US-0363124P.

2002WO-US017674

11-MAR-2002; 29-MAY-2002; 06-JUN-2002;

03-JUL-2002;

2002US-0386782P 2002US-0393796P

2002US-0399348P 2002US-0406784P. 2002US-0408378P.

09-SEP-2002; 2002US-0409293P.

05-SEP-2002; 7-NOV-2002;

29-AUG-2002;

15-JAN-2003; 2003US-0440129P

2002US-00306747

Human VEGFR1 short interfering nucleic acid (siNA) SEQ ID NO:585

(first entry)

12-FEB-2004

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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothalial growth factor receptor (VEGFF) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, nephrotropic and gynaecological, antiatritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating downregulating) the expression of VEGFF genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 8 A; 1 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 158; 207pp; English.
                                                                                                                                                                                                                                                                20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0353124P.
29-MAY-2002; 2002WG-US017674.
06-JUN-2002; 2002US-0386782P.
03-JUL-2002; 2002US-0393796P.
29-JUL-2002; 2002US-039348P.
                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
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27-NOV-2002; 2002US-00306747.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-679876/64.
                                                                                                                                            WO2003070910-A2
                                                                                                         Homo sapiens.
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                                                                                                                                                                                    28-AUG-2003
                                                                                    Synthetic
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGER) gene. Also described: (1) a sinA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of engine-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 3 A; 7 C; 1 G; 0 T; 8 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor gene.
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Mcswiggen J, Beigelman L, (RIBO-) RIBOZYME PHARM INC

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Gaps

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0.4%; Score 15.8; DB 1; Length 19; 99.5%; Pred. No. 1.2e+03; ve 0; Mismatches 2; Indels

89.5%;

Conservative

Best Local Similarity Matches 17; Conserv

Query Match

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Gaps

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2; Indels

3; Mismatches

1294 GTGAAGATGCTGAAAGACG 1312 GUGAAAAUGCUGAAAGAGG 19

14; Conservative

Matches

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ADF36296 standard; RNA; 19 BP

ADF36296 RESULT 1035 ADF36296/C ID ADF36290 XX AC ADF36290

Score 15.8; DB 1; Length 19; Pred. No. 1.2e+03;

0.4%;

Query Match Best Local Similarity

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1 GUGGCCGUGAAAUGCUGA 19
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0386782P.
                                                                              0.4%;
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2002US-00306747.
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                                                                                                                                                                                                                             ADF37398 standard; RNA; 19
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                                                                                                                                                                                                                                                                                 (first entry)
                                                                                         Local Similarity 68.4 es 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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11-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAY-2002;
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                                                                              Query Match
                                                                                                                                                                                                RESULT 1037
ADF37398
                                                                                            Best Local
Matches
883333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a double-stranded short interfering nucleic acid (stNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA; with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating to (downregulating) a vide range of angiogenesis-associated conditions, useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, chung screening, target identification and validation, genetic
                                                                                                                                                                                                double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; short interfering nucleic acid; siNA; downregulation; cantidial growth factor receptor; VEGFR; antianglogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; alabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                        Human VEGFR3 short interfering nucleic acid (siNA) SEQ ID NO:1650
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1294 GTGAAGATGCTGAAAGACG 1312
                                                                                                                                                                                                                                                                                                                                                                                                                                            20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-035124P.
29-MAY-2002; 2002WS-03517574.
06-JUN-2002; 2002US-0386782P.
03-JUL-2002; 2002US-0393796P.
29-JUL-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
05-SEP-2002; 2002US-0408378P.
                                                                                           BP
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27-NOV-2002; 2002US-00306747.
                GIGAAATGCIGAAAGAGG
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                                                                                         ADF37361 standard; RNA; 19
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                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                         13
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    (e.g. o
in the
                                                                                                                                                                                                                                                                                                                                                            Gaps
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engineering, studying gene function, and also for gene mapping single-nucleotide polymorphisms). The present sequence is used exemplification of the present invention.
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                                                                                                                                                                                                                                                       Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
4; Mismatches 2; Indels
                                                                                                                                                                         Sequence 19 BP; 5 A; 3 C; 7 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          1288 GTAGCCGTGAAGATGCTGA 1306
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RESULT 1039
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                                                                             (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, barticularly cancers, diabetic retinopathy, macular degeneration, neovascular y claubetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  short interfering nucleic acid; sing, downcaulation; vascular endothelial growth factor receptor, VBGPR, antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotroppic, gynaecological, angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma;
  express siNA; and (5) single-stranded siNA with similar properties.
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                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                     The sinhs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating
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0
                                                                                                                                                                                                                                                                                                                                                                    Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 4 C; 7 G; 0 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        double-stranded short interfering nucleic acid;
                                                                                                                                                                                                                                                                                                                                                                                                               7; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              1807 TGGTCCTTTGGGGTCCTGC 1825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 UGGUCCUUUGGGGUGCUUC 19
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29-MAY-2002; 2002WG-US011674.
06-UUN-2002; 2002US-0386782P.
29-UUL-2002; 2002US-0393796P.
29-AUG-2002; 2002US-039348P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADF37608 standard; RNA; 19 BP
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09-SEP-2002; 2002US-0409293P.
04-NOV-2002; 2002US-00287949.
                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%;
Best Local Similarity 52.6%;
Matches 10; Conservative 7
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15-JAN-2003; 2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1038
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGER) gene. Also described: (1) a siNA that downregulates the VEGF gene, (2) kits for in vitro or in vivo of siNA, (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antianglogenic, cytostatic, antidabetic, ophthalmological antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiotibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, angineering, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the
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and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 7 C; 3 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid;
                                                                                                       Example 3; SEQ ID NO 1897; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         exemplification of the present invention.
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2002US-0409293P.
2002US-00287949.
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                                          factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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09-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma;
                                                                                                                        New double-stranded interfering nucleic acid, useful e.g. for treatment
                                                                                                                                       and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                              Example 3; SEQ ID NO 1935; 207pp; English.
                                                                    Beigelman L, Pavco P;
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27-NOV-2002; 2002US-00306747
15-JAN-2003; 2003US-0440129P
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                                        (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                       factor receptor gene.
                                                                                             WPI; 2003-679876/64.
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                                                                    Mcswiggen J,
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antianglogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynascological activities. The siNA are useful for andulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, barcularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 1688; 207pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1825 CTCTGGGAGATCTTCACGC 1843
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20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
29-MAY-2002; 2002WO-US017674.
06-UUN-2002; 2002US-0386782P.
03-UUL-2002; 2002US-0393796P.
29-UUL-2002; 2002US-039378P.
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                                                                                                                                                                                                                           29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                                                                                                                    09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                             27-NOV-2002; 2002US-00306747
                                                                                                                                                                                                                                                                                                                                                                                                                   L5-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mcswiggen J, Beigelman L,
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Best Local Similarity 63.2
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        factor receptor gene.
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The present invention describes a double-stranded short interfering nucleic acid (sinA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a sinA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of sinA; (3) conjugates and/or complexes of sinA; (4) vectors that express sinA, and (5) single-stranded sinA with similar properties. The sinAs have antialangiogenic, cytostatic, nephrotropic and gynaecological activities. The sinA are useful for modulating ophthalmological, antiatrhritic, antipsoriatic, nephrotropic and gynaecological activities. The sinA are useful for modulating consequiating the expression of VEGFR genes. The sinA are potentially particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, arbititis, psoriasis, endometriosis, angiofibroma, and plycystic kidney disease. The sinA may also be useful for diagnosis, dung screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of single-incledide polymorphisms). The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double-stranded interfering nucleic acid, useful e.g. for treatment diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.8; DB 1; Length 19;
89.5%; Pred. No. 1.2e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 8 A; 7 C; 4 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beigelman L, Pavco P;
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                                                                                                                                                                                                                                        06-UIN-2002; 2002US-0386782P.
03-UIL-2002; 2002US-0393148P.
29-UIL-2002; 2002US-0393148P.
29-XUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-04092949.
27-NOV-2002; 2002US-00306747.
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                                                                                                                                                20-FEB-2003; 2003WO-US005022
                                                                                                                                                                                      2002US-0358580P
                                                                                                                                                                                                      11-MAR-2002; 2002US-0363124P.
29-MAY-2002; 2002WO-US017674
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Best Local Similarity
                                                                       WO2003070910-A2
               Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J,
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ID ADF478

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The invention relates to short interfering nucleic acids (siNA) which comparises to superassion of the human Myc or Myb genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense creations, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNAs), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, candor complexes of siNA; and vectors that express siNA. The siNAs are andor complexes of siNA; and vectors that express siNA, in siNAs are capanisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be consed for treatment of a variety of conditions. They may be consed for treatment of a variety of conditions. They may be consed for treatment of a variety of conditions and so useful for treatment of a variety of conditions and so useful for drug screening, diagnosis, therapeutic target identification and the present sequence represents the upper strand of a human Myc-targeted double-stranded siNA, which is identical to the Myc transcript target
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                                     Human; Myc; Myb; cancer; proliferative disease; restenosis; polycystic kidney disease; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cytostatic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of Myc or Myb genes
Human Myc transcript target sequence/siNA upper strand, SEQ ID 17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; Page 126; 161pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0409293P.
; 2002US-0418655P.
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                                                                                                                                                                                                                                                                                                                                                                                                                             2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                                 nephrotropic; ss.
                                                                                                                                                                                                                                                                                WO2003070917-A2.
                                                                                                                                                                                                                                           Homo sapiens.
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11-MAR-2002;
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Gaps ö

vivlemore401-10.rng

Length 19;

Query Match

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Human; Myc; Myb; cancer; proliferative disease; restenosis; polycystic kidney disease; RNA interference; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cytostatic; vasotropic;
                                                                                                                                                                                                                                        Human Myc siNA lower strand, SEQ ID 135.
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06-JUN-2002; 2002US-0386782P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-0409293P.

15-GCT-2002; 2002US-0418655P.
                                                               ADF47998 standard; RNA; 19 BP
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                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nephrotropic; ss
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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of Myc or Myb genes Example 7; Page 126; 161pp; English.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human Myc or Myb genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense cligonaclectide. The siNAs include short an antisense oligonaclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express sinAs are used to modulate expression of the Myc or Myb genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and creatment conflexed, diseases, such as restenosis and polycystic kidney disease. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene conflexed, services sinas and seconds of the Myc or Myb genes in cells (function, and gene mapping (e.g., of single nucleotide polymorphisms).

Sequence 19 BP; 4 A; 3 C; 10 G; 0 T; 2 U; 0 Other;

double-stranded siNA.

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                                                                                                                                                                                                                                                                                                        88; 8iNA; human; BCL2; short interfering nucleic acid; RNA interference; cytostatic; immunosuppressive; virucide; anti-HIV; cancer; autoimmune disease; viral infection; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel short interfering nucleic acid (sink) that downregulates expression of the BCL2 gene by RNA interference. A sain of the invention has cytostatic, immunosuppressive, virucide, and anti-HIV activity. The sink are useful for modulation (inhibition) of expression or activity of BCL2 by RNA interference. sink are used to modulate expression of BCL2 genes, in cells, tissue explants or organisms, e.g. for treating cancer, autoimmune diseases and viral infections (including by HIV) but also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (e.g. of sing nucleotide polymorphisms). The sequences shown in ADF49273-ADF50143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer or autoimmune disease, downregulates expression of the BCL2 gene.
                                     Gaps
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                                 Indels
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0.4%; Score 15.8; DB 1;
78.9%; Pred. No. 1.2e+03;
ive 2; Mismatches 2;
                                                                                                                                                                                                                                                                             Human BCL2 siNA lower sequence SEQ ID NO:130.
                                2; Mismatches
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                                                                   1950 GATCATGCGGGAGTGCTGG 1968
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2002US-0363124P.
2002US-0386722P.
2002US-0406784P.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409233P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                ADF49402 standard; RNA; 19
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                  Local Similarity 78.5
nes 15; Conservative
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                    Best Loca
Matches
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Human IGF-1R transcript target sequence/siNA upper strand, SEQ ID NO:192.

12-FEB-2004 (first entry)

ADF31527;

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; micro-RNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; autoimmune disease; transplant rejection; cytostatic; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human; insulin-like growth factor 1 receptor; IGF-IR; target sequence; ss.

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The invention relates to a novel short interfering nucleic acid (siNA) that downregulates expression of the BCL2 gene by RNA interference. A soll A of the invention has cytostetic, immunosuppressive, virucide, and anti-HIV activity. The siNA are useful for modulation (inhibition) of expression or activity of BCL2 by RNA interference. siNA are used to modulate expression of BCL2 genes, in cells, tissue explants or organisms, e.g. for treating cancer, autoimmune diseases and viral infections (including by HIV) but also for drug screening, diagnosis, target identification and validation, genetic engineering, phammacogenomics, studying gene function and gene mapping (e.g. of single nucleotide polymorphisms). The sequences shown in ADF49273-ADF50143
                                                                                                                                                                                                                     ss; siNA; human; BCL2; short interfering nucleic acid; RNA interference; cytostatic; immunosuppressive; virucide; anti-HIV; cancer; autoimmune disease; viral infection; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer or autoimmune disease, downregulates expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.8; DB 1; Length 19; 36.8%; Pred. No. 1.2e+03; ive 10; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 8 G; 0 T; 11 U; 0 Other;
                                                                                                                                                                                     Human BCL2 siNA lower sequence SEQ ID NO:544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 544; 148pp; English.
19 TIGITGIGIGIGIGI 1
                                                                                  ADF49816 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2002; 2002US-0358580P
11-MAR-2002; 2002US-0363124P
06-JUN-2002; 2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUL-2002; 2002US-0396905P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                  18-FEB-2003; 2003WO-US004908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 36.55
Best Local 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-712622/67.
                                                                                                                                                                                                                                                                                                                             WO2003070969-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the BCL2 gene.
                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                     12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                28-AUG-2003
                                                                                                                    ADF49816;
                                                RESULT 1045
ADF49816
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human insulin-like growth factor 1 creceptor (IGFT-IR) gene by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They comprise sense and may be double or single stranded. They further comprise sense and matisense regions, or alternatively are sesembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be chemically synthesised, expressed from a decoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits or sinA; and vectors that express siNA. The siNAs are used to modulate expression of the IGF-IR gene in cells, tissue explants or organisms of sinA; and vectors that express siNA. The siNAs are used to modulate expression of the IGF-IR gene in cells, tissue explants or organisms or treatment of a variety of conditions. They may be used for treating cancer and other proliferative diseases (e.g., restenosis and polycystic kidney diseases), inflammatory and/or allergic diseases, autoimmune diseases and transplant rejection. The siNAs are also useful for drug crecening, diagnosis, therapeutic target identification and validation, and gene companies, therapeutic target identification and validation, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence crepresents the upper strand of a human IGF-IR transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 192; 147pp; English.
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Gaps

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2312 TIGGICIGIGIGIGIGI 2330

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ADF31527 standard; RNA; 19 BP

RESULT 1046 ADF31527 ID ADF31527

Sequence 19 BP; 4 A; 8 C; 4 G; 0 T; 3 U; 0 Other;

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the insulin-like growth

factor-1 receptor gene.

Chowrira B;

Mcswiggen J, Beigelman L, (RIBO-) RIBOZYME PHARM INC

WPI; 2003-721691/68.

11-MAR-2002; 2002US-0363124P. 06-UIN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408178P.

15-JAN-2003; 2003US-0440129P

20-FEB-2003; 2003WO-US005044.

20-FEB-2002;

WO2003070911-A2. Homo sapiens.

28-AUG-2003.

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SINA.
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      8$888888888$
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; micro-RNA; short hairpin RNA; shRNA; expression modulation; gene therapy; phag screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; autorimmune disease; transplant rejection; cytostatic; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human; insulin-like growth factor 1 receptor; IGF-1R; ss.
                                                                   Gaps
                                                                   ö
Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human IGF-1R siNA lower strand, SEQ ID NO:469.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 469; 147pp; English.
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                                                                                                                             1617 CCACAGGGACCTGGCTGCC 1635
                                                                                                                                                                                       1 ccacacacaccuudcuece 19
                                                                                                                                                                                                                                                                                                                                                 BP
      0.4%;
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06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409293P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JAN-2003; 2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                 ADF31804 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      factor-1 receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-721691/68.
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                                   Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003070911-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J,
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                                                                                                                                                                                                                                                                                                                                                                                                               ADF31804;
      Query Match
                                                                                                                                                                                                                                                                              RESULT 1047
APP31804/C
AX
ADP31804/C
ADP31804/C
AX
ADP31804/C
ADP3
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treatment of a variety of conditions. They may be used for treating cancer and other proliferative diseases (e.g., restenosis and polycystic kidney disease), inflammatory and/or allergic diseases, autoimmune diseases and transplant rejection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human IGF-IR-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-vabl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint
                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
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                                                                                                                                                                                                                         Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                          Sequence 19 BP; 3 A; 4 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7; SEQ ID NO 840; 197pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ABL1-targeted siRNA - SEQ ID 840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cluster region-Abelson (BCR-ABL) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chowrira
                                                                                                                                                                                                                                                                                                        1617 CCACAGGGACCTGGCTGCC 1635
                                                                                                                                                                                                                                                                                                                                             19 ccacadadaccrrdcrdcc 1
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2002US-0408378P.
2002US-0409293P.
2003US-0439922P.
2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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2002US-0363124P.
2002US-0386782P.
2002US-0404039P.
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                                                                                                                                                                                                                               0.4%;
                                                                                                                                                                                                                                                   89.58;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF84546 standard; RNA; 19
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                                                                                                                                                                                                                             Query Match 0.4
Best Local Similarity 89.5
Matches 17; Conservative
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05-SEP-2002;
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11-MAR-2002;
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ς΄

miRNA;

short interfering RNA; siRNA; double stranded RNA; micro-RNA; miRNA, short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HERR2; EGFR2; neu; erbB2; c-erb-B-2; ss.

short interfering nucleic acid; siNA;

RNA interference;

Human HER2 (EGFR2) siNA lower strand, SEQ ID NO:403.

20-MAY-2004 (first entry)

BP.

ADL79238 standard; RNA; 19

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ADL79238;

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The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-v-abl Abelson murtine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                         short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
                                                                                                     ö
                                                                  0.4%; Score 15.8; DB 1; Length 19; 63.2%; Pred. No. 1.2e+03; Live 5; Mismatches 2; Indels
                                   Sequence 19 BP; 2 A; 5 C; 6 G; 0 T; 6 U; 0 Other;
the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 7; SEQ ID NO 1159; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                          Human ABL1-targeted siRNA - SEQ ID 1159.
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                                                                                                                                   3688 TICTIGGGGCCCAGTGCAT 3706
                                                                                                                                                        1 UUCUCGGGGUCCAGUGCAU 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
15-AUG-2002; 2002US-0404039P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-040878P.
09-SEP-2002; 2002US-0408378P.
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                                                                                                                                                                                                                                                     ADF84865 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                  Local Similarity 63.2
nes 12, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003070972-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-AUG-2003
                                                                    Query Match
                                                                                                                                                                                                                                                                                         ADF84865;
                                                                                                                                                                                                                      RESULT 1049
                                                                                                                                                                                                                                       ADF84865
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29-MAY-2002; 2002WO-US016840. 06-JUN-2002; 2002US-00163552.

11-MAR-2002; 29-MAY-2002;

20-FEB-2003; 2003WO-US005045

WO2003070912-A2. Homo sapiens.

28-AUG-2003

06-JUN-2002; 2002US-0386782P. 03-JUL-2002; 2002US-0393924P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P.

09-SEP-2002; 2002US-0409293P. 19-SEP-2002; 2002US-00251117.

15-JAN-2003; 2003US-0440129P

(RIBO-) RIBOZYME PHARM INC.

2002US-00277494

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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                             Jamison S;
                                                                                                                                              Fosnaugh K,
                                                                                                                                                                            Example 3; SEQ ID NO 403; 171pp; English.
                                                                                                                                              Mcswiggen J, Pavco P, Beigelman L,
                                                                                                                                                                     factor receptor gene.
                                                                                                                                                    WPI; 2003-697612/66.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain decoxyribonucleotides, and can be chemically synthesised, can contain decoxyribonucleotides, and can be chemically synthesised, can contain decoxyribonucleotides, and can be chemically synthesised, and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of EGFR genes in cells, tissue explants or organisms (eg., by ex vivo gene therappy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The
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Gaps

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Query Match 0.4%; Score 15.8; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 1.2e+03; Matches 17; Conservative 0; Mismatches 2; Indels

Sequence 19 BP; 6 A; 6 C; 5 G; 0 T; 2 U; 0 Other;

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Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL79242;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loc
Matches
      88888888888888888888888888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human HER2 (EGFR2) transcript target sequence/siNA upper strand, SEQ:158.
siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a HER2 (EGFR2)-targeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HBR1, HBR2 HBR3 and HBR4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New short interfering nucleic acid, useful e.g. for treatment and. diagnosis of cancer, downregulates expression of the epidermal growth factor receptor gene.
                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; dragnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; erbB2; c-erb-B-2; target sequence; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jamison S;
                                                                                                                                                                         Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beigelman L, Fosnaugh K,
                                                                                                                              Sequence 19 BP; 5 A; 6 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; SEQ ID NO 158; 171pp; English.
                                                                                                                                                                                                                                                           1672 ATCGCAGACTTCGGGCTGG 1690
                                                                                                                                                                                                                                                                                      19 ATTACAGACTTCGGGCTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002; 2002WO-US016840.
06-UJN-2002; 2002US-00163552.
06-UJN-2002; 2002US-038582.
03-UJL-2002; 2002US-0383934P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                              ADL78993 standard; RNA; 19 BP
                                                                                                                                                                         0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002US-0358580P.
2002US-0363124P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2003; 2003WO-US005045
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                     17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pavco P,
                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-697612/66.
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11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-SEP-2002;
21-OCT-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL78993;
                                                                                                                                                                                                                                                                                                                                                                      RESULT 1051
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                        ADL/8993

110

ADL/78993

20

XX

ADL/2

XX

ADL/2

XX

ADL/2

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cc regions, or alternatively are assembled from a sense oligonucleotide and an antiense oligonucleotide. Specifically, the sinks include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also cand/or complexes of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of EGFR genes in cells, tissue explants or corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants or creating a wide range of cancers such as breast and ovarian cancer. The citreating a wide range of cancers such as breast and ovarian cancer. The cathoristication and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide collable stranded siNA, which is identical to the upper strand of a characting a wide and double-stranded siNA, which is identical to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; agne function analysis; gene mapping; cancer; cytoetatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; erbB2; c-erb-B-2; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.8; DB 1; Length 19; 73.7%; Pred. No. 1.2e+03; ive 3; Mismatches 2; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human HER2 (EGFR2) siNA lower strand, SEQ ID NO:407.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J, Pavco P, Beigelman L, Fosnaugh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 4 A; 6 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the HER2 transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1744 CCCGTGAAGTGGATGGCGC 1762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002US-0363124P
2002US-0363124P
2002US-0316540
2002US-0386782P
2002US-040674P
2002US-0406784P
2002US-0406293P
2002US-0402293P
2002US-0402293P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 73.7%;
les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADL79242 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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Sequence 19 BP; 3 A; 6 C; 6 G; 0 T; 4 U; 0 Other;
                                   Example 3; SEQ ID NO 407; 171pp; English.
                                                                                                                                                                                                                              1744 CCCGTGAAGTGGATGGCGC 1762
                                                                                                                                                                                                                                       19 CCCATCAAGTGGATGGCGC 1
                                                                                                                                                                                                                                                                        ADL78989 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
29-MAY-2002; 2002US-0US016840.
06-UUN-2002; 2002US-00163552.
06-UUN-2002; 2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2003; 2003WO-US005045.
                                                                                                                                                                                                                                                                                                20-MAY-2004 (first entry)
                       factor receptor gene.
WPI; 2003-697612/66.
                                                                                                                                                                                                                                                                                                                                                                                     WO2003070912-A2
                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                28-AUG-2003.
                                                                                                                                                                                                                                                                                    ADL78989;
                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                             RESULT 1053
ADL78989
                                                                                                                                                                                                                   Matches
8
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convergulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA increference. The siNAs may or may not comprise sense and antisense of double or single stranded. They further comprise sense and antisense of oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be enemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNAs are used to modulate expression of EGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants or creating a wide range of cancers such as breast and ovarian cancer. The treatment of a variety of conditions. They may be used for treating a mide range of cancers such as breast and ovarian cancer. The creating are also useful for drug screening, diagnosis, therapeutic target complication, and gene mapping (e.g., of single nucleotide collymorphisms). The present sequence represents the upper strand of a content of the present sequence represents the upper strand of a content of the content of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human; T-cell associated disease; Vbeta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                   Mcswiggen J, Pavco P, Beigelman L, Fosnaugh K, Jamison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 4 A; 4 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 154; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human Vbeta gene repeat sequence #23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the HER2 transcript target sequence.
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1 AUUACAGACUUCGGGCUGG 19
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                                2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                19-SEP-2002; 2002US-00251117.
21-OCT-2002; 2002US-00277494.
15-JAN-2003; 2003US-0440129P.
2002US-0393924P
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                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.4
Best Local Similarity 68.4
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-697612/66.
                                29-AUG-2002;
05-SEP-2002;
                                                                                                   09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH70233;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH70233,
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                                                                                                                                                                                                                                                         The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor downregulate expression of one or more human epidermal growth factor receptors (including HER1, HER2 HER3 and HER3 by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (micRNA) and short can contain deoxyribonucleotides, and can be chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Can contain deoxyribonucleotides, and can be chemically synthesised. Can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of BGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants or creating a wide range of cancers such as breast and ovarian cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target conduction, and gene mapping (e.g., of single nucleotide collaboration), genetic empires in the lower strand of a chemically such as breast and corporation of ethicity (e.g., of single nucleotide conducts). The present sequence represents the lower strand of a chemical such as breast such as breast and corporation of a stranded sinA.
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                                                                       New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA interference; short interfering nucleic acid; siNh; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HERZ; EGFRZ; neu; exbBZ; c-erb-B-Z; target sequence; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             / Match 0.4%; Score 15.8; DB 1; Length 19; Local Similarity 89.5%; Pred. No. 1.2e+03; hes 17; Conservative 0; Mismatches 2; Indels
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Gaps

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degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; aultiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; doodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection;

vivlemore401-10.rng

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enigma; TROP2; mitosin; CU-EC-1; cytostatic; solid cancer; RT-PCR;
primer; ss.
                                                                                                                                                                                                                                   Shimada H, Hiwasa T,
                                                                                                                                                                                                                                                                          WPI; 2004-248065/23
                                                                                               WO2004018518-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2004018679-A1.
                                                                                                                                                                                                                                                Takiguchi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens.
                                                       Homo sapiens
                                                                                                                          04-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-MAR-2004
                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADL71963;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo
                 ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     셤
                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Wheta gene, WetaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases including autoimmune disease, degenerative nervous system diseases, infectious diseases including autoimmune disease, whypersensitivity diseases, infectious diseases and neoplastic disease, Autoimmune diseases include Addison's disease, atrophic gastritis. Degenerative nervous system diseases include multiple arrophic gastritis. Degenerative nervous system diseases include multiple arrophic gastritis as contact with allergens that lead to allergies. Type II hypersensitivities such as those present in callergies. Type II hypersensitivities such as those present in codpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by viruses such as HUY, fungal infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by Wycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                              Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                           filaria, bacterial infection, Mycobacterium, neoplastic disease,
lymphoproliferative disease, leukaemia, lymphoma, cancer, brain cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Solid-cancer antigen; FIR; CENP-A; HOOK2; myomegalin; MKRN1; KIAA1545;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
               fungal infection; Candida; parasitic infection; schistosome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match 0.4%; Score 15.8; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 1.2e+03; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 10 A; 0 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 427; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA amplifying RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2834 ATATATATATACATATAT 2852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19 ATATATATATATATAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADL72177 standard; DNA; 19 BP.
                                                                                                                                                                  99US-00263959
                                                                                                                                                                                           94US-00309335.
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                                                                                                                                                                                                                                                                                                      WPI; 2004-059052/06
                                                                                                                                                                                                                                                                             Hood LE, Rowen L;
                                                       breast cancer; ds
                                                                                                                                                                                                                                   (HOOD/) HOOD L E. (ROWE/) ROWEN L.
                                                                                                            US2002150891-A1
                                                                                                                                                                                           19-SEP-1994;
19-SEP-1995;
                                                                                  Homo sapiens
                                                                                                                                                                  05-MAR-1999;
                                                                                                                                     17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                           Vbeta gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human FIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADL72177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1055
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The invention relates to human solid-cancer antigen peptides and encoding polynucleotides. The antigen peptides are selected from FIR, CENP-A, HOOK2, myomegalin, WRNI, KIAA1545, enigma, TROP2, mitosin, CU-EC-1. Cytostatic. The antigen peptides and their polynucleotides are applicable in early diagnosis and useful in treating solid cancer. With these 10 antigen peptides and their encoded polynucleotides, early diagnosis of solid cancer can be achieved accurately and quickly. The present sequence represents a RT-PCR primer for amplifying the DNA encoding a human solid-cancer antigen peptide FIR.
                                                                                                                                                                                                                                                                                                                                                                  Human solid cancer antigen peptides and polynucleotides useful in early
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FIR; CENP-A; cancer antigen peptide; far-upstream binding protein interacting receptor; centromere-specific protein A; cancer; human; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.8; DB 1; Length 19;
Pred. No. 1.2e+03;
                                                                                                                                                                                               Nomura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 2 A; 4 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                               Tomonaga T, Matsushita K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human FIR DNA amplifying PCR primer, SEQ ID 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; SEQ ID NO 23; 136pp; Japanese.
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                                                                                                                             (NISC-) JAPAN SCI & TECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis and treating solid cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1189 CTGACCCTGGGCAAGCCCC 1207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADL71963 standard; DNA; 19 BP.
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21-APR-2003; 2003WO-JP005046.
                                                       23-AUG-2002; 2002JP-00244249.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                           Ochiai T;
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criteria. Also claimed are methods for gene-silencing, developing an sirah algorithm for selecting siRNA, selecting an siRNA with improved unctionality, selecting siRNA, selecting an siRNA with improved effective at silencing byperfunctional siRNA, an siRNA with improved siRNA. The siRNA molecule comprises a sequence substantially similar to a sequence consisting of GGGAGAUGAGGAGGA, GAAGUCAUCAUCAUGAGG, CAUGCGCCUCGGAGANA, AGAUGAGGAGGACAAU, GAAGUCACCCGAGAUA, AGAUGAGGAGGAGCAC, The siRNA molecule comprises a sense strand and an anti-sense strand. The siRNA molecule comprises a hairpin. The siRNA molecule comprises between 18 and 30 base pairs. The kit comprises at least two siRNA, comprising a first optimised siRNA for generating a gene silencing reagent. The present sequence is sired in the exemplification of the invention. The sequence is shown in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= a //note= at least one (and preferably all) of the backbone subunits are composed of N-acetyl N-(2-aminoethyl)glycine peptide residues, the nucleobase being attached covalently to the acetyl group and the peptide linkage being formed by condensation of the glycine carboxy group of one residue with the amino group of the 2-aminoethyl moiety in the next residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptide nucleic acid oligomers specific for protein kinase C isozyme(s) - useful as anti:sense molecules for treating PKC mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide nucleic acid; PNA; PKC-alpha; protein kinase C; ss; cell proliferation; cell differentiation; isozyme; antisense; triple helix; cancer; psoriasis; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PNA oligomer targetting coding region of PKC-zeta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the specification as DNA, but described as siRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 7 A; 0 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.8; DB 1;
Pred. No. 1.2e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1294 GTGAAGATGCTGAAAGACG 1312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GTGAAGATGTTGAAGATG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94WO-US008465.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 89.5'
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-082040/11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JUL-1993;
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18-OCT-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ97929;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1058
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                 THE STATE OF THE S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                            The invention relates to novel polynucleotides encoding all or part of cancer antigen peptide FIR (far-upstream binding protein interacting receptor) or peptides derived from it and polynucleotides encoding all or part of cancer antigen peptide CENP-A (centromere-specific protein A) or a peptide derived from it. The polynucleotides or primers, probes and substances containing them are use in a method for cancer diagnosis and especially diagnosis of colorectal cancer. The present sequence represents a PCR primer for amplifying the human FIR DNA.
                                                                                                                                                                           Cancer antigen polypeptides, nucleic acids encoding them and antibodies recognising them for specific and sensitive diagnosis of colon and rectal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Selecting siRNA by selecting an siRNA molecule of 19-25 nucleoside bases by selecting a target gene and measuring the functionality of the nucleotide sequences that are complementary to a stretch of nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel method for selecting siRNA (short interfering RNA) comprising selecting an siRNA molecule of 19-25 nucleoside bases by selecting a target gene and measuring the functionality of sequences of 19-25 nucleotides in length that are substantially complementary to a stretch of nucleotides of the target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene silencing; Bcl-2; optimised; short interfering RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ô
                                                 Nomura F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.8; DB 1; Length 19; 89.5%; Pred. No. 1.2e+03; ive 0; Mismatches 2; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stephen S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-FGFR2 siRNA related DNA sequence SEQ ID NO:720.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 2 A; 4 C; 11 G; 2 T; 0 U; 0 Other;
                                             Ochiai T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 William M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 12; SEQ ID NO 720; 199pp; English.
                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 8; 323pp; Japanese.
                                                 Matsushita K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Devin L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1189 CTGACCCTGGGCAAGCCCC 1207
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10-SEP-2003; 2003US-0502050P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 89.5
les 17; Conservative
                                             Tomonaga T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Angela R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (DHAR-) DHARMACON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-420527/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         interference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2004045543-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-JUN-2004
                                             Shimada H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      siRNA;
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Synthetic.

BB;

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RESULT 1057

ADQ61018

ADQ61018

Query Match

Best Loca Matches

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Gaps

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Example 15; Page 36; 125pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                          AAT30427;
                                                                                                                                                                                                                                                                                                                                                                    RESULT 1060
                                                                                                                                                                                                                                                                                                                                                                                   AAT30427,
                                                                                                                                                                                                                                                                                                                                                                                               셤
                                                                                                                                                                                                                                                                                                  ઠે
                                              New peptide nucleic acid (PNA) oligomers are provided which (a) consist of naturally occurring nucleobases covalently bound to a polyamide backbone and (b) hybridise to the translation intiation AUG region, coding region, 5' untranslated region (5' UTR) or 3' untranslated region (3' UTR) of PKC-alpha or its isoforms. The PNAs can be used to target RNA and simple stranded DNA (sBNNA) to produce antisense-type gene regulation moieties. They inhibit expression of PKC-alpha and its isoforms (including beta, gamma, delta, epsilon, zeta and eta) and so are useful cor treating and diagnosing cell proliferation and differentiation processes such as neoplastic, hyperproliferation and differentiation or processes such as neoplastic, hyperproliferative and inflammatory diseases. PNA oligomers have high affinity for complementary single tranded DNA. They are also able to form triple helices in which a first PNA strand binds with RNA or sBNNA and a second PNA strand binds with the resulting double helix or with the first PNA strand binds with the submitted of an an analysic of supplication charge and are water soluble, which facilitates cellular uptake. Further, since they contain amides of non-biological amino acids, they are biostable and resistant to enzymatic degradation by proteases. The present sequence targets the coding region of PKC-zeta. (Updated on
                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense; protein kinase C; alpha; PKC; beta; gamma; eta; epsilon; zeta; modulation; expression; isozyme; hybridise; 5' UTR; human; y untranslated region; translation initiation site; detection; phosphorothioate linkage; 2'-0-methyl modification; 2'-0-propyl modification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligo:nucleotide(s) hybridisable with Protein Kinase C mRNA or gene -also novel PKC-alpha 3'-UTR sequence, useful for diagnosis and treatment of hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PKC-zeta coding region antisense oligo, ISIS #9010.
                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 1 A; 8 C; 3 G; 8 T; 0 U; 0 Other;
disease, e.g. cancer, psoriasis and inflammation
                         Claim 31; Page 270; 287pp; English
                                                                                                                                                                                                                                                                                                  25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                            323 CTCCCTCCATCTCCTGGCT 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dean NM;
                                                                                                                                                                                                                                                                                                                                                                                                                                      crrccrccarcrrcreecr 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94US-00199779.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93US-00089996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ84246 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                   17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bennett CF, Boggs RT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-066911/09.
                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9502069-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-JUL-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
21-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ84246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1059
                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ84246
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The sequences given in AAQ84244-51 are oligos which are antisense to the protein kinase C-zeta (PKC-zeta) cDNA. These antisense molecules may be used in modulating the expression of this particular isozyme of PKC. These oligomers have a *inhibition of PKC of <70%. The oligos of the invention preferably hybridise with the 5'- or 3'-untranslated regions of the invention preferably hybridise with the 5'- or 3'-untranslated regions of the invention preferably hybridise with the 6'- or 3'-untranslated regions of the invention preferably hybridise with the 6'- or 3'-untranslated regions of the end oligos may be used in the detection of the human PKC genes and for treatment of animals with conditions associated with PKC, esp.

These oligos may be used in the detection of the human PKC genes and for treatment of animals with conditions associated with PKC, esp.

Appropriate linkage and/or at least one of the nucleotides comprises a modification of the sugar, esp. a 2'- or methylor a modification of the sugar, esp. a 2'- or methylor a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the probes. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor segment. The present sequence is an example of a compound SSR primer. The method represents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             obsition of the sugar, esp. a 2'- O-methyl or a (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; compound; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compound simple sequence repeat primer (CA)4.5(TA)7.5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 8 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DUPO ) DU PONT DE NEMOURS & CO E I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   323 CTCCCTCCATCTCCTGGCT 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 crrccrccarcrrcrescr 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95WO-US015150.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-00346456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT30427 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2'-0-propyl modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JAN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.4
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Morgante M, Vogel JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-277795/28.
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junB gene antisense oligonucleotide JunB-T-11.
                                                                                                                                                                       Brysch W;
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                                                                                                             97EP-00101531
                                                                                                                               97EP-00101531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV73071 standard; DNA; 20
                                                                                                                                                                                           WPI; 1998-400910/35.
                                                                                                                                                                       Schlingensiepen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
es 17; Conserv
                                                                                                             31-JAN-1997;
                                                                                                                               11-JAN-1997;
                                                  Homo sapiens
                                                                    EP856579-A1
                                                                                         05-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-FEB-1999
                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV73071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a serum albumin-growth hormone fusion protein useful to treat growth hormone related diseases such as Down's syndrome. This polynucleotide sequence represents the DNA of a fusion region of the pHGH56 and pHGH57 growth hormone protein plasmids of the invention
                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                              Serum albumin-growth hormone fusion protein - useful to treat growth hormone related diseases, e.g. Down's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                              Fusion region of pHGH56 and pHGH57 growth hormone protein plasmids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
0
                                                           ö
                                                                                                                                                                                                                                  hormone fusion protein; growth hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 15.8; DB 1; Length 20; 73.7%; Pred. No. 1.3e+03; tive 4; Mismatches 1; Indels
                                     Query Match 0.4%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.3e+03; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 0 A; 0 C; 16 G; 0 T; 0 U; 4 Other;
                   Sequence 20 BP; 10 A; 7 C; 0 G; 3 T; 0 U; 0 Other;
 genetic trait marking and germplasm comparisons
                                                                               2338 TGTGTGTGTGTGCACAT 2356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2920 GGGCGGGCGTGGGGGGC 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 11; 21pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                   (DELZ ) DELTA BIOTECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 rererererererarar 2
                                                                                                                                                   AAK99571 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                      95GB-00026733.
96WO-GB003164.
                                                                                                                                                                                                                                             Down's syndrome; chimeric; ds
                                                                                                                                                                                                                                                                                                                                    98KR-00704914.
                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 73.7
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV48702 standard; DNA;
                                                                                                                                                                                                                                  Serum albumin-growth
                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-363680/33
                                                                                                                                                                                           06-AUG-2002
                                                                                                                                                                                                                                                                                             KR99076789-A.
                                                                                                                                                                                                                                                                                                                                    25-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                      30-DEC-1995;
19-DEC-1996;
                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                        Ballance DJ;
                                                                                                                                                                                                                                                                                                               15-OCT-1999.
                                                                                                                                                                                                                                                                         Chimeric.
                                                                                                                                                                       AAK99571;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV48702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1062
AAV48702
ID AAV48702
XX
AC AAV48702
AC DT 15-OCT-1
XX
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AAV48564-708 represent antisense oligonucleotides directed against the jumb and jumb genes. Of these, only oligonucleotides AAV4855-614
resulted in effective downregulation of negative growth control by Jumb or Jumb, while AAV48615-708 had little effect. The oligonucleotides exemplify the invention. The specification describes oligonucleotides that can each form three hydrogen bonds to cytosine; do not contain four consecutive rytosines; do not contain two sequences of three consecutive mucleotides able to form three H-bonds each to four consecutive cytosines; and the ratio between residues able to form two H-bonds each (2R) or three such bonds (3R) is given by 2R/3R = 0.33-0.72. The oligonucleotides are used to modulate expression of genes, particularly the genes for p53, ErbB-2, jumb, jumb, TGF-beta 1 or beta 2 to control consideration of primary cell cultures (e.g. bone marrow stem, liver or kidney cells, osteoclasts, osteoblasts and/or keratinocytes). The consecution also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (coligonucleotides can also be used to analyse function of proteins (coligonucleotides can also be used to analyse function of proteins (coligonucleotides can also be used to analyse function of proteins (coligonucleotides can also be used to analyse function of proteins (coligonucleotides can also be used to analyse coligonucleotides can also be used to analyse coligonucleotides can also be asset to analyse coligonucleotides can also be asset coligonucleotices (coligonucleotides can also be asset coligonucleotides can are coligonucleotides can also be asset coligonucleotices of col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             altering their expression or activity) and therapeutically, e.g. in cases of cancer or (targeting TGF) for stimulating the immune system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in
junB; junD; antisense oligonucleotide; modulate; gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
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                                                                                                                                                                                                  Ras oncogene; probe; point mutation; detection; cancer; ss.
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necrosis factor alpha (TNP-alpha) 5'-flanking region. The present invention describes genetic variations involved in diseases associated with TNF-alpha which are screened for by detecting the presence or absence of one or more of the following mutations in the 5'-flanking region of the human TNF-alpha gene: (1) T in place of C at position -857; (2) A instead of C at -853; and (3) C instead of T at -1031 [positions 373; 367 and 199 respectively of the sequence given in AAX18575). The mutations increase TNF-alpha production in the cells involved. A suitable method for detection of these mutations is allele-specific oligonucleotide hybridisation (ASOH). TNF-alpha spentic variations associated with TNF alpha-implicated diseases such as juvenile rheumatoid arthritis, insulin-dependent diabetes or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; tumour necrosis factor alpha; TNP-alpha; promoter; diagnosis; juvenile rheumatoid arthritis; chronic rheumatoid arthritis; insulin-dependent diabetes; insulin-independent diabetes; pCR primer; ss.
Diagnosis of genetic variation associated with diseases involving TNF alpha - such as rheumatoid arthritis or diabetes, by detecting mutation at positions 857, 863 or 1031 of the TNF alpha gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Diagnosis of genetic variation associated with diseases involving TNF alpha - such as juvenile rheumatoid arthritis, by detecting mutation at positions 857, 863 or 1031 of the TNF alpha gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human tumour necrosis factor alpha 5'-flanking region PCR sense primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                       sequence represents a PCR primer for the human tumour
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.8; DB 1; Length 20; Pred. No. 1.3e+03;
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                                                                    Disclosure; Page 5; 34pp; Japanese.
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97JP-00173900.
97WO-JP004304.
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                                                                                                                                                                                                                                                                                                                                    insulin-independent diabetes
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Best Local Similarity 89.5
Matches 17; Conservative
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26-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                  AAV73026-V73071 are probes used to detect a single-base mutation in a human ras oncogene. These probes comprise 12-43 nucleotides of formula 5'-B-Q-D-3', Q=3 nucleotides complementary to the mutated codon, and B and D each = 0-20 nucleotides complementary to the ras sequences flanking the mutated codon. The probes are useful for detecting cancers associated with point mutations
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                                                                                                                    85US-00758104.
87US-00081490.
92US-00873352.
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                                                                                                                                                                                                           (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                             Van Der Eb AJ;
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04-AUG-1987;
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with TNF-alpha which are screened for by detecting the presence or absence of one or more of the following mutations in the 5'-flanking region of the human TNF-alpha gene: (1) T in place of C at position -857; (2) A instead of C at -863; and (3) C instead of T at -1031 (positions mutations in crease TNF-alpha production in the cells involved. A suitable mutations increase TNF-alpha production in the cells involved. A suitable method for detection of these mutations is allele-specific or oligonucleotide hybridisation (ASOM) TNF-alpha genetic variations associated with TNF alpha-implicated diseases such as juvenile rheumatoid arthritis, chronic rheumatoid arthritis, insulin-dependent diabetes or
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Pred. No. 1.3e+03;
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Local Similarity 89.5%;
les 17; Conservative
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The invention relates to the isolation of a nucleic acid (AAA11284) coding for a transmembrane protein TRPC7 (AAY92944) which is expressed in brain and is homologous to transient receptor potential (TRP) protein. This suggests that the TRPC7 protein may have a calcium channel function. The genomic sequence has been shown to contain 31 introns. This sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding transmembrane protein TRPC7 expressed in brain and homologous to transient receptor potential protein useful in the of treatment of associated diseases such as periodic psychosis.
ALIS gene from a sample for comparison to normal samples in the in vitro diagnosis regime. This may also be performed by amplifying XIIS CDNA from the mRNA in the sample. Antibodies to XIIS may be used to detect XIIS in a biological sample or can be administered to patients to prevent or treat the above disorders. They may also be used to patients to prevent or treat the above neurological sample. XIIS may also be used to purify XIIS from a biological sample. XIIS may also be administered to patients to prevent or treat the above neurological disorders. In addition XIIS may be used as a marker of neuronal cells at an early stage of development; its discovery increases understanding of both the neuronal movement which leads to development of the cortical region of the brain and of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Transmembrane protein; TRPC7; brain; transient receptor potential; TRP; calcium channel function; human; gene therapy; periodic psychosis;
                                                                                                                                                                                                                                                                                                                       Gaps
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Pred. No. 1.3e+03;
0; Mismatches 2;
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/number= 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA11330 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimizu N, Nagamine K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .10
/*tag=
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                                                                                                                                                                                                                                                                                                       Local Similarity
les 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200029571-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mutation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .2-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA11330;
                                                                                                                                                                                                                                                                                      Query Match
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intron
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1067
                                                                                                                                                                                                                                                                                                                             Matches
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DNA fragmentation; microsatellite DNA; DNA marker;

(first entry)

99WO-JP003551.

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Isolation of satellite seguences from genomic DNA for use as DNA markers comprises isolating a library with high homogeneity by DNA fragmentation.
                                                                                                                                                                                                                                                                                                                           (NORQ ) JAPAN MIN AGRIC FORESTRY & FISHERIES.
                                                                                                                        H. discus derived sequence #17.
                                         AAZ98499 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                      rakahashi H, Sekino M;
                                                                                                                                               Satellite sequence;
Haliotis discus; ss.
                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-224692/19.
                                                                                                                                                                                         Haliotis discus.
                                                                                                                                                                                                                    WO200011156-A1.
                                                                                                                                                                                                                                                                        01-JUL-1999;
                                                                                             19-JUN-2000
                                                                                                                                                                                                                                             02-MAR-2000
                                                                   AAZ98499;
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                           AAZ98499
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represents an exon/intron junction from the genomic TRPC7 sequence. The DNA and protein can be used in the diagnosis and treatment of disorders associated with TRPC7, especially the screening, monitoring and treatment (by gene therapy) of periodic psychosis, which appears to be associated with mutations in the TRPC7 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method which comprises making a high-density gene chip, specifically for making high-density micro-array of oligonucleotide probes. An oligonucleotide probe selecting process to seek preferentially length variable and coverage variable probes is provided to ensure identical cross melting temperature of probes to the maximum limit, and this can make the cross control of gene chip relatively simple and raise the relability of the gene chip results. The process proposes a specific probe selection method for and non-specific stres and a probe overall arrangement scheme. AAA79738 to AAA80201 represent oligonucleotide probe sequences which are used in
                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus; HBV; Hepatitis A virus; HAV; probe; detection;
mutation; high-density gene chip; ss.
                                                                                                                                       ö
                                                                                                       Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; rive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                         Hepatitis B virus related oligonucleotide probe #186.
                                                                                 Sequence 20 BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 6 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High-density gene chip making process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   examples from the present invention
                                                                                                                                                               1566 TGCCTACCAGGTGGCCCGG 1584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Fig 15; 19pp; Chinese.
                                                                                                                                                                              TGCCTTCCAGGTGGCCCAG 19
                                                                                                                                                                                                                                                        AAA79923 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99CN-00114460.
                                                                                                           0.4%;
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                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                      17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYDO-) UNIV DONGNAN.
                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis B virus.
                                                                                                                                                                                                                                                                                                             20-NOV-2000
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The invention provides a novel method for isolation of satellite sequences from genomic DNA that comprises fragmentation of the DNA by a method which is not dependent on base sequences, then selection of the satellite sequences from the obtained genomic library of high homogeneity. The method is useful for the isolation of microsatellite DNA sequences which can be used as DNA markers. The new method markedly improves the efficiency of isolation of satellite sequences in comparison to prior art methods which are reliant on base sequences. Sequences AAZS9483-514 represent sequences from Haliotis discus, used in the method
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antiinflammatory, 2'-methoxyethyl wing; 2'-MOE wing; infection; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
0.4%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 1.3e+03;
Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human fra-1 mRNA antisense oligonucleotide ISIS 109023.
                                                                                                                                                                                                                                                                                                                                                              Seguence 20 BP; 6 A; 10 C; 4 G; 0 T; 0 U; 0 Other;
Example 5; Page 14; 35pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2329 GIGIGCGIGIGIGIGIG 2347
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Gaps

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AGGAGCTCTTCAAGCTGCT 1895

Conservative

17; 1877

Best Loc Matches

Local Similarity

Query Match

19 AGGAGCTCTTCTAACTGCT 1

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inflammatory, autoimmune and heart disease.
                                                                                                                                                                                                                                                                                                                                      inhibiting inflammation or apoptosis
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                                                                                  Example 5; Page 53; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                   43 GGGCCCCAGCGGCTGCAGG 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 grececadedecrecade 2
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            WPI; 2000-664982/64.
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Best Local Similarity
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                                    Antisense
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                                                                                                                                                                             Novel antisense compound 8-30 nucleobases in length targeted to human fra -1 and which specifically hybridizes with and inhibits the expression of human fra-1, useful for modulating the expression of fra-1 in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotide; p38 mitogen activated protein kinase; MAPK; antirknematic; antiarthritic; immunosuppressive; cardiant; heart disease; antiinflammatory; autoimmune disease; rheumatoid arthritis; apoptosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                The present sequence is one of a large number of antisense oligonucleotides which are targeted to nucleic acids encoding fra-1. The sequences may be oligodecoxyribonucleotides or chimeric oligonucleotides containing a central gap region consisting of ten 2'-decoxynucleotides, which is flanked on both sides by 2'-methoxyethyl (2'-WDE) wings. The oligonucleotides have a phosphorothioate backbone and the cytidine residues in the 2'-MDE wings are 5-methylcytidines. The fra-1 antisense oligonucleotides are useful for inhibiting the expression of fra-1 in human cells or tissues. They can be used for diagnostics, therapeutics, prophylaxis and as research reagents and in kits. Use of the antisense compounds may also be useful prophylactically, e.g. to prevent or delay
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaarde WA, Nero PS, Mckay R, Popoff I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Murine p38beta antisense oligonucleotide SEQ ID 65.
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3.A; 8 C; 9 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                   inflammation or tumour formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1484 GGCGCCCCCGGGCCTGGA 1502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGCGGCCCAGGGCCCGGA 20
                                                                                                                                                                                                                            Claim 3; Col 41; 38pp; English.
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                                                         99US-00418641
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                                                                                 99US-00418641
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Les 17; Conservative
                                                                                                                               Cowsert LM;
                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                        WPI; 2000-601552/57
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                                                         15-OCT-1999;
                                                                                 15-OCT-1999;
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           US6124133-A.
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                                  26-SEP-2000
                                                                                                                                Taylor JK,
                                                                                                                                                                                                                                                                                                                                                                                      infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monia BP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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translational translated region, translational start site, translational translational start site, translational termination region or 3'-untranslated region of a nucleic acid encoding a p38 mitogen activated protein kinase (MAPK), where the antisense oligonuclectides inhibit the expression of MAPK. Sequences AAC79501 represent human p38alpha MAPK and p38beta MAPK cDNA sequences. AAC79501 a AAC79500 and AAC79532 - AAC79570 represent human p38alpha antisense oligonuclectides, while AAC79502 - AAC79521 and AAC79571 - AAC79580 represent human p38alpha antisense oligonuclectides while AAC79536 solated from rat tissue. Murine p38beta MAPK CDNA is represented in AAC79533 and antisense oligonuclectides AAC79523 - AAC79536 isolated from rat tissue. Murine p38beta MAPK CDNA is represented in AAC79537 and antisense oligonuclectides have antifficumatic; antisting and inhibiting the expression of p38 MAPK in cligonuclectides are useful for inhibiting the expression of p38 MAPK in cells or tissues. The oligonuclectides are used for treating an animal with diseases such as inflammatory or autoimmune diseases e.g. rheumatoid arthibities, or heart disease. The oligonuclectides are used for treating an animal with this titles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                     This invention relates to antisense compounds 8-30 nucleobases in length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for mechanically sensitive potassium channel gene fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
Antisense compound targeted to p38 mitogen activated protein kinase inhibits protein kinase and is useful for diagnosing and treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 10 C; 6 G; 2 T; 0 U; 0 Other;
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Gaps

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14-FEB-2001; 2001WO-US004732
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                                                                                                                                          AAD15530 standard; DNA; 20
                                                                                                                                                                                                                                      Bollon AP, Gray DM,
                                                                                                                                                                                                                      (BOLL/) BOLLON A P. (GRAY/) GRAY D M.
                                                                                                                                                                                                                                              WPI; 2001-529916/58
                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                               JU-SEOG L.
                                                                                 neurodegeneration
                                                                                                                                                                                        WO200161030-A2
                                                                                                                                                         15-NOV-2001
                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                23-AUG-2001
                                                                                                                                                  AAD15530;
                                                                                                                                                                                                                               (JUSE/)
                                                                                                                                   RESULT 1073
                                                                                                       Matches
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sequences, as well as antisense oligonucleotides capable of binding DNA. The antisense and antigene libraries are useful for preparing therapeutic agents for the treatment of genetic disease. The present DNA sequence is human C-Raf protein target DNA related to the invention. Note: The present sequence is shown as DNA in the specification; however, in vivo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method for selecting optimal subsequence antisense targets. The method involves preparing an antisense oligonucleotide capable of inhibiting mRNA expression of target mRNA sequences, as well as antisense oligonucleotides capable of binding DNA. The antisense and antisens libraries are useful for preparing therapeutic agents for the treatment of genetic disease. The present DNA sequence is phosphorothioate antisense oligonucleotide which is targetted to human C-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Selecting optimal subsequence antisense targets for inhibition of mexpression of target mRNA for the therapeutic treatment of genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; C-Raf protein; genetic disease; therapeutic; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                                                   0.4%; Score 15.8; DB 1; Length 20; ilarity 89.5%; Pred. No. 1.3e+03; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human C-Raf gene targetted antisense oligonucleotide #1
                                                                                                                                                            Sequence 20 BP; 4 A; 3 C; 12 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Phosphorothicate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                this target sequence would be mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 19; 87pp; English
                                                                                                                                                                                                                                                                                             843 GCTGCCAGCCGAGGAG 861
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/mod_base=
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-529916/58
                                                                                                                                                                                                                          Local Similarity
nes 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (JUSE/) JU-SEOG L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200161030-A2
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-AUG-2001
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                                                                                                                                                                                                     Query Match
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(GRAY/) (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Raf gene
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Matches
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                                                                                                                                                                   PCR primers AAH78639-40 were used to amplify a gene fragment of the human mechanically sensitive potassium channel gene. The channel is activated by polyumsaturated fatty acids (particularly arachidonic acid (AA)) and by riluzole. The polypeptide is designated human TWICK-related AA-activated potassium channel (hTRAAK). The hTRAAK gene is located on chromosome 11q13. hTRAAK is involved in regulation of neuronal and muscle excitation, cardiac rhythm and secretion of hormones. Cells that express hTRAAK, designated to screen for modulators of hTRAAK activity. Such modulators are potentially useful for prevention or treatment, in humans and animals, of: cardiac and/or vascular disease; nervous system disocated with anomalous hormone secretion or muscle diseases associated with anomalous hormone secretion or muscle diseases and retinal diseases. Typical examples are epilepsy, cardiac arrhythmia and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; C-Raf protein; genetic disease; antisense target; therapeutic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Selecting optimal subsequence antisense targets for inhibition of mexpression of target mRNA for the therapeutic treatment of genetic disease.
                                         New mechanically sensitive potassium channel, useful for treating cardiovascular diseases and in drug screening, is activated by polyunsaturated fatty acids.
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Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                              Disclosure; Page 15; 37pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human C-Raf protein target DNA #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1869 CCCTGTGGAGGAGCTCTTC 1887
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCAGTGGAGGAGCCCTTC 20
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WPI; 2001-590037/66
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The present invention describes a method for promoting the inhibition of the expression of a protein comprising employing a mammalian RNase H polypeptide so that cleavage of an RNA strand of an oligonucleotide-RNA complex duplex occurs. Also described is a compound 8 to 50 nucleobases in length targeted to the nucleic acid encoding the human RNase HII polypeptide, where the compound specifically hybridises with and inhibits the expression of a human RNase HII polypeptide. The compound, which is an antisense oligonucleotide, is useful for inhibiting the expression of a human RNase HII polypeptide in cells or tissues, as well as for a human RNase HII polypeptide in cells or tissues, as well as for treating an animal with a disease or condition associated with a human CR RNAse HII polypeptide. The method is useful for inhibiting the expression of a protein, particularly for reducing cellular RNA via antisense technology. The present sequence represents a human RNASE HII antisense coligonucleotide, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of a mammalian, particularly human, RNase H, for treating an animal with a disease or condition associated with a human RNase H, for inhibiting the expression of a protein, or for reducing cellular RNA via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unexplained recurrent pregnancy loss; immunologic reproductive failure; URPL; interleukin-loeta; IL-lbeta; human; tumour necrosis factor a; promoter; PCR; primer; ss.
                                                                                                                                                                             /mod_base= OTHER /nod_base= OTHER /note= "2'-O-methoxyethyl gapmer with an 8 nucleotide deoxy gap and a phosphorothioate backbone; cytosine residues are 5-methyl cytosines"
            RNase H; antisense technology; inhibition; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 12 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human TNFA promoter amplifying PCR primer #1.
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89.5%; Pred. No. ...
                                                                                                                 Location/Qualifiers
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Best Local Similarity 89.5
Matches 17; Conservative
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                                      phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     technology
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modified_base
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                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense
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MAKAKEKAKAKAKA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an antisense compound 8 to 30 bases in length targeted to the 5' untranslated (UTR) region, the coding region or the 3' UTR region human Integrin-linked kinase (hIK). The antisense oligonuclectides are useful for inhibiting the expression of human hIK in human cells or tissues, in vitro. The oligonuclectides can be utilized for diagnostics, therapeutics for the treatment of diseases associated with the expression of hIK, prophylaxis e.g. to prevent or delay infection, inflammation or tumor formation and as research reagent
                                                                                                                                                                                                                                                                                                                                                                                   Antisense; integrin-linked kinase; hIK; infection; tumour; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel antisense compounds capable of modulating expression of human Integrin-linked kinase, useful for diagnosis, prophylaxis and treatment of diseases, e.g. tumors, associated with expression of the kinase.
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                                        Gaps
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                                        Indels
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. 1.3e+03;
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                 Pred. No. 1.36
0; Mismatches
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                                                                               843 GCTGCCAGCCGAGGAG 861
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                                                                                                                       19 ścreccaecedaceaea 1
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                 89.5%;
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                     Best Local Similarity 89.5
Matches 17; Conservative
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es 17; Conservative
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RESULT 1076

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ABQ75387/ ID ABQ7 XX AC ABQ7 AC ABQ7 DT 06-N XX XX

Matches

nucleic

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therapy. The KTOMI nucleic acids may be used to diagnose, treat or monitor a disease caused by altered expression of human KTOMI.

Compositions comprising the nucleic acids, proteins or antibodies may be used to treat subjects having defects in KTOMI which can manifest as cancer of the kidney, as well as a disorder of liver, bone marrow, brain, heart, lung, kidney, colon, skeletal muscle, testis, uterus and placenta function. The sequence represents a PCR primer used in the invention to
                                                                                                                                                                                                                                                                                                                                                                                                New human kidney tumor overexpressed membrane (KTOM1) protein and nucleic acids encoding the protein, useful for treating subjects having defects in KTOM1 which can manifest as cancer of the kidney, or as a disorder of e.g., liver or bone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated nucleic acid encoding human KTOM1 (kidney tumour overexpressed membrane) protein. The protein of the invention has cytostatic activity. The nucleotide may have a use in gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse; antiinflammatory; gene therapy; ileitis; DST; ss; primer; real-time validation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 100; 418pp; English.
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                30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-US000669.
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                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-479509/51.
                                                                                                                                                                                                                                                                (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200231114-A2
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                                                                                                                                                                                                                                                                                                            Zhang J;
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                                                                                                                                                                                                                                                                                                                                                                                                  Evaluating risk of unexplained recurrent pregnancy loss in a subject, by testing presence of a variant in interleukin-1 beta promoter region and/or in CD46 gene intron 1 region in a sample obtained from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method for evaluating and treating risk of unexplained recurrent pregnancy loss (URPL) in a subject suspected of having immunologic reproductive failure. The method involves testing a sample obtained from the subject for the presence of a variant in the human interleukin-lbeta (IL-lbeta) promoter region, and/or a variant in the CD46 gene intron 1 region, where the presence of the variant, indicates an elevated risk of developing recurrent pregnancy loss. The present sequence is human tumour necrosis factor a (TNFA) promoter amplifying PCR primer used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; KTOM1a; KIOM1; kidney tumour overexpressed membrane; cytostatic;
gene therapy; cancer; kidney; liver; bone marrow; brain; heart; lung;
kidney; colon; skeletal muscle; testis; uterus; placenta; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         r match 0.4%; Score 15.8; DB 1; Length 20; Local Similarity 89.5%; Pred. No. 1.3e+03; eB 17; Conservative 0; Mismarchae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 2 C; 9 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human KTOMla control RT_PCR primer hNAC-1F3.
                                                                                                                                                                                                                                                                                                            Hill JA, Wang ZC, Anderson DJ, Yunis EJ;
                                                                                                                                                                                                                                            (BGHM ) BRIGHAM & WOMENS HOSPITAL INC. (DAND ) DANA FARBER CANCER INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 21; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2333 GCGTGTGTGTGTGTGTG 2351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GCTTGTGTGTGTGTCTG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
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27-SEP-2000; 2000US-023539P.
04-OCT-2000; 2000GB-00024263.
30-JAN-2001; 2001WO-US000661.
                                                                                                                                                      12-SEP-2001; 2001WO-US028465
                                                                                                                                                                                                12-SEP-2000; 2000US-0231785P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-2001; 2001WO-US029656
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                                                              WO200222877-A2
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                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-AUG-2002
                                                                                                            21-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-MAR-2002
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RESULT 1078

ABQ65252

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04-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABS57369;
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                                                                                                                                                                           The invention relates to a novel isolated nucleic acid molecule comprising a polynucleotide having one of 90 polynucleotide sequences, given in the specification. The polynucleotides of the invention have antiinflammatory activity, and may have a use in gene therapy. The polynucleotide or a polypeptide encoded by it is used for preventing, treating, modulating or ameliorating a medical condition such as ileits. The polypeptide or polynucleotide is also useful for manufacturing a medicament for treating ileitis. The sequence represents a real-time validation primer for the DNA sequence obtained from one of the mouse
                                                                    New isolated nucleic acid molecules that are associated with ileitis, for preventing, treating, modulating and diagnosing ileitis in a mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      neuroprotective; cerebroprotective; nootropic; cytostatic; cardiant; nephrotropic; osteopathic; antiallergic; antiarteriosclerotic; anti-microbial; diabetes; infection; dementia; cytochrome P; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Transgenic animals having drug metabolism enzyme genes, useful in testing fetotoxicity and/or teratogenicity and applicable to drug development for diseases including diabetes, infections and dementia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a recombinant gene that comprises, a gene encoding human P450 or its variant, the human EFlalpha promoter, chick beta globin insulator sequence or a part of it, and the SV40 polyA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Transgenic animal; drug; fetotoxicity; teratogenicity; antidiabetic;
 Hilbush BS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ô
                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kato M;
 Hasel KW,
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Teranishi Y, Ishida M,
 Baum PR,
                                                                                                                                            Disclosure; Page 227; 273pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 27; 60pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2687 AGGCTTTCCCACTTCCCAC 2705
 Dubose RF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 AGGCTGTCCCACATCCCAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABQ81623 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          17; Conservative
                                                                                                                                                                                                                                                                                                                                                  clones of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
 Sims JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CYP2E1 sense primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENC-) GENCOM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-674938/72.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-AUG-2002
Viney JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABQ81623;
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                                                                                                            subject
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ABQ81623/c
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attached signal. The activity of the gene of the invention may be described as, antidiabetic, neuroprotective, cerebroprotective, nootropic, osteogethic, artialergic, nootropic, categorathic, antiallergic, antiarteriosclerctic and anti-microbial. The intention of this invention is to provide a transgenic animal. The animal is useful in testing fectoxicity and/or teratogenicity, and is applicable to drug development for diseases including diabetes, infections and dementia. The current sequence represents a primer designated OYP2E1 sense primer, which is used in an example from the invention in the amplification of cytochrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human cancer cell growth suppressing protein PP6781 cDNA, PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ø
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; cancer cell growth suppression; cancer; PP6781; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human protein with cancer cell growth suppressing function and polynucleotide encoding it, for treating diseases, such as, cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 8 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.8; DB 1;
Pred. No. 1.3e+03;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                     P from total RNA from human liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  842 TGCTGCCAGCCGAGGA 860
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-OCT-2000; 2000CN-00127101.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS57369 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 89.5'
Matches 17; Conservative
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10-MAY-2003 (first entry)

ABZ74966;

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New antisense compound, useful for preparing a composition for diagnosing, treating or preventing inflammatory diseases, e.g. rheumatoid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "nucleotides 1-5 & 16-20 are 2'-methoxyethoxy (MOE) nucleotides, nucleotides 1-4 & 16-19 are linked via phosphodiester linkages, nucleotides 6-15 are 2'-deoxy- nucleotides, nucleotides 5-16 are linked via phosphorothicate linkages, all C nucleotides are 5-methyl cytosines"
                                                                                                                                                                                                                                                                                                                                                               p38 mitogen-activated protein kinase; p38 MAPK; phosphorothioate; antisense; antiarthritic; antiinflammatory; kinase inhibitor; mouse; inflammatory disease; rheumatoid arthritis; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel antisense compound, which is 8-30
                                                                                                                                                                                                                                                                                                Murine p38-alpha MAPK antisense oligonucleotide ISIS NO 100802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.8; DB 1; Length 20; Pred. No. 1.38+03; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 10 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mckay R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Col.27-28; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                             ABX78139 standard; DNA; 20 BP
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                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-089122/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                          16-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               arthritis
                                                                                                                                                    ABX78139;
RESULT 1082
ABX78139/C
ID ABX78139/C
XX AC ABX78139/C
XX IG-APR-:
XX DT 16-APR-:
XX MUSH MUSH
XX MUSH MUSH
XX MUSH MUSH
XX MUS
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Sequences ABZ74952-ABZ74991 represent antisense oligonucleotides targeted to the human p70 S6 kinase gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human p70 S6 kinase RNA, and were analysed for their effect on mRNA levels by quantitative real-time PCR. p70 S6 kinase (also known as SK6, p70/p85 s6 kinase, p70/p85 ribosomal S6 kinase and pp70s6k) is a serine-threonine kinase responsible for the phosphorylation of the ribosomal S6 protein, which in turn stimulates protein synthesis. p70 S6 kinase function is essential for cell cycle progression, and has also been implicated in the regulation of the immune response. p70 S6 kinase is itself activated via phosphorylation, a process influenced by upstream signalling cascades and by hyperinsulinaemia, and may play a role in the progression of colon cancer and in the development of lipotoxic of the progression of disorders
                                                                                                                                                                                                                                                                                                                                                                                                     /note= "Phosphorothioate linkages. When bases 1-5 and 16-20 are not 2'-methoxyethyl (2'-MOE) nucleotides, all cytosines in the oligonucleotide are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel antisense compound which is targeted to nucleic acid encoding p70 S6 kinase, and inhibits expression of p70 S6 kinase, useful for treating a condition associated with p70 S6 kinase, e.g. cancer.
                                                                                                                              Human; p70 S6 kinase; SK6; p70/p85 s6 kinase; pp70s6k; pp70/p85 ribosomal S6 kinase; serine-threonine kinase; ribosomal S6 kinase; serine-threonine kinase; ribosomal S6 protein phosphorylation; protein synthesis; cell cycle progression; immune response; signalling cascade; cancer progression; liptocxic disorder; obsativ; metabolic disorder; hyperproliferative disorder; cancer; cytostatic; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and obesity. The oligonucleotides of the invention are useful for the prevention and treatment of conditions associated with p70 S6 kinase, such as hyperproliferative disorders such as cancer, and metabolic disorders. They are also useful in research and diagnostics for modulating the expression of p70 S6 kinase
                                                                                            Human p70 S6 kinase phosphorothioate antisense oligo, SEQ ID NO:24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2'-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Optionally 2'-methoxyethyl (2'-MOE;
All 2' MOE cytosines are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 73; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2002; 2002WO-US023123.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-AUG-2001; 2001US-00920677.
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                                                                                                                                                                                                                                                                                                                                                                *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003012032-A2
                                                                                                                                                                                                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                        Homo sapiens
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Gaps

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0.48; 89.58;

Query Match

Local Similarity 89.5 es 17; Conservative

ABZ74966 standard; DNA; 20 BP

RESULT 1083 ABZ74966/c ID ABZ7496

ss; PCV-2; PCR; primer; RFLP; restriction fragment length polymorphism; postweaning multisystemic wasting syndrome; PWWS; anti-viral; porcine circovirus infection.

Porcine circovirus type 2.

US2003096377-A1.

22-MAY-2003.

(VIRG) VIRGINIA TECH INTELLECTUAL PROPERTIES.

WPI; 2003-606420/57.

Meng X, Fenaux M;

27-JUN-2002; 2002US-00184191. 28-JUN-2001; 2001US-0301707P.

Porcine circovirus PCV-2 sequencing/PCR primer CV 3.

(first entry)

06-NOV-2003

ADA09089;

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BP.

ADA09089 standard; DNA; 20

ADA09089/

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The invention relates to the isolated polynucleotides of a repeated tract within intron 1 of the zinc finger protein 9. The isolated polynucleotides comprise nucleotides 1-14468, 14444-22400, 17501-17701 and optionally a repeat tract, 17858-18062 and optionally a repeat tract, of a 22400 base pair sequence given in the specification, or its complements, or at least about 15 consecutive nucleotides from 16701-17701 or 17858-18862 of the 22400 by sequence. The polynucleotides of the invention are useful in identifying individuals at risk for developing myotonic dystrophy type 2 (DM2). This polynucleotide sequence represents a PCR primer of the human zinc finger protein 9 of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated polynucleotide for determining whether an individual has, is at risk, or is not at risk for developing myotonic dystrophy type 2, comprises a repeat tract within intron 1 of the zinc finger protein 9
                                                                                                                                                                                                                                                                                                                                       tract; intron 1; zinc finger protein 9; myotonic dystrophy type 2;
                                                             Gaps
                                                             ô
                            Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
Sequence 20 BP; 2 A; 12 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                      Zinc finger protein 9 DNA PCR primer SEQ ID No 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 1 A; 2 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 21; 66pp; English.
                                                                                           853 GAGGAGGAGCTGGTGGAGG 871
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Liquori C;
                                                                                                                                                                                                          ABT16306 standard; DNA; 20 BP
                                                                                                                          GAGGATGAGCTGGAGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-MAY-2001; 2001US-0290365P.
29-JUN-2001; 2001US-0302022P.
13-NOV-2001; 2001US-0337831P.
                            Query Match
Best Local Similarity 89.5%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-MAY-2002; 2002WO-US014837
                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MINU ) UNIV MINNESOTA.
                                                                                                                                                                                                                                                                                                                                                      DM2; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RANUM L P W.
DAY J W.
LIQUORI C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ranum LPW, Day JW,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genomic sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                    WO200292763-A2
                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                        20-MAR-2003
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                                                                                                                          20
                                                                                                                                                                                                                                          ABT16306;
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(DAYJ/)
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                                                                                                                                                                                                                                                                                                                                       Repeat
                                                                                                                                                                          RESULT 1084
                                                                                                                                                                                         ABT16306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to detecting and differentiating porcine circovirus (PCV) infections, comprises extracting nucleic acid from a biological asmaple taken from a pig, amplifying a fragment from the extracted nucleic acid, digesting the amplified fragment with a restriction earyme, forming a restriction fragment length polymorphism (RFLP) pattern from an undigested or digested fragment, and detecting the presence or absence of a PCV isolate. Also included are an oligonuclecide primer for differentiating PCV infections, comprising a fully defined MCV1 or MCV2 mucleotide sequences (ADA09085 and ADA09086) and an assay kit for detecting and differentiating PCV infections, comprising the oligonucleotide primers and a restriction enzyme. The methods and compositions of the present invention are useful for diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treating porcine circovirus infection (especially postweaning multisystemic wasting syndrome, PMWS, in pigs). The method of the present invention, as compared to prior art, is more rapid, sensitive, easy to perform and specific for the porcine circovirus and its associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is a PCR/sequencing primer for performing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting and differentiating porcine circovirus infections utilizing differential polymerase chain reaction-restriction fragment length polymorphism assay using a restriction enzyme and primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.8; DB 1; Length 20; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2420 CTGCTGTGCAACGGTCTCC 2438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 9; 21pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB25743 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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ADB25743
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ADB25743;

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Gaps

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Score 15.8; DB 1; Length 20; Pred. No. 1.3e+03; 0; Mismatches 2; Indels

0.4%;

Local Similarity 89.5 es 17; Conservative

Query Match

Best Loca Matches

2325 GTGTGTGTGCGTGTGTG 2343

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crererecentriciere 19

Mouse connective tissue growth factor antisense oligo DNA (SeqID 136) antisense; mouse; murine; ss; connective tissue growth factor; CTGF; chromosome 6q23.1; ctgrofact; fibroblast inducible secreted protein;

ВР.

ADB25743 standard; DNA; 20

RESULT 1087

20-NOV-2003

ADB25743;

/mod_base= OTHER /note= "OTHER= phosphorothioate backbone, where 1-5 and 16-20 are 2' methoxyethyl nucleotides. All cytidines are 5-methylcytidines"

09-DEC-2002; 2002WO-US038618. 10-DEC-2001; 2001US-00006191.

WO2003053340-A2

03-JUL-2003

(ISIS-) ISIS PHARM INC.

Gaarde WA, Watt AT;

Location/Qualifiers

antiarteriosclerotic

ds snw

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/*tag=

modified base

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fisp-12; NOVŽ; insulvative binding protein-related protein 2; IGFBP-rP2; IGBBP-8; Hcs24; ecogenin; acute lymphoblastic leukaemia; gene therapy; hyperproliferative disorder; cancer; pulmonary fibrosis; renal fibrosis; scleroderma; atherosclerosis; cytostatic; dermatological;
                                                                                                                                                                                                                                  /mod_base= OTHER /nosphorothioate backbone, where 1-5 and /note= "OTHER= phosphorothioate backbone, where 1-5 and 16-20 are 2' methoxyethyl nucleotides. All cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides for modulating connective tissue growth factor expression, particularly useful for treating cancers (e.g. breast or prostate cancer), pulmonary or renal fibrosis, scleroderma or atherosclerosis.
                                   Mouse connective tissue growth factor antisense oligo DNA (SeqID 136)
                                                         mouse; murine; ss; connective tissue growth factor; CTGF; 6q23.1; ctgrofact; fibroblast inducible secreted protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 11 A; 1 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                       5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Page 89; 139pp; English.
                                                                                                                                                                                                                                                                                                                                                                       10-DEC-2001; 2001US-00006191.
                                                                                                                                                                                                                                                                                                                                               09-DEC-2002; 2002WO-US038618.
           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                 antiarteriosclerotic
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaarde WA, Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-559091/52.
                                                                                                                                                                                                                                                                                               WO2003053340-A2.
                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                        03-JUL-2003
                                                           antisense;
chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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This invention relates to novel methods for modulating the expression of connective tissue growth factor (CTGF) by antisense oligonucleotides.

CTGF has been mapped to human chromosome region 6423.1, and is also known as ctgrofact, fibroblast inducible secreted protein. 2, 1618p-12, 100V2, insulin-like growth factor binding protein related protein. 2, 1GFBP-12, IGFBP-18, Hcs24 and ecogenin. It is known to stimulate DNA synthesis and promote chemotaxis of fibroblasts, however, it is also upregulated in accute lymphoblastic leukaemia and in tumour or endothelial cells associated with the vasculature. Accordingly, antisense oligonucleotides that inhibit the expression of CTGF in cells or tissues can be used in gene therapy to treat various conditions including hyperproliferative disorders (particularly cancer, e.g. breast, prostate or renal cancer), pulmonary fibrosis, renal fibrosis, scleroderma and atherosclerosis. As
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      such, the present invention describes these antisense oligos as having cytostatic, dermatological and antiarteriosclerotic activities. This oligonucleotide sequence is a chimeric phosphorothioate antisense oligo with 2' MOE wings and a deoxy gap, which is used to inhibit expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides for modulating connective tissue growth factor expression, particularly useful for treating cancers (e.g. breast or prostate cancer), pulmonary or renal fibrosis, scleroderma or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Page 89; 139pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-559091/52.
This invention relates to novel methods for modulating the expression of connective tissue growth factor (CTGF) by antisense oligonucleotides.

CTGF has been mapped to human chromosome region 623:1, and is also known as ctgrofact, fibroblast inducible secreted protein, fisp-12, NOV2, insulin-like growth factor binding protein-related protein 2, IGFBP-rP2, IGFBP-8, Hose24 and ecogenin. It is known to stimulate DNA synthesis and IGFBP-8, Hose24 and ecogenin. It is known to stimulate DNA synthesis and coute lymphoblastic leukaemia and in tumour or endothelial cells associated with the vasculature. Accordingly, antisense oligonuclectides that inhibit the expression of CTGF in cells or tissues can be used in gene therapy to treat various conditions including hyperproliferative disorders (particularly cancer, e.g. breast, prostate or renal cancer), cub, the present invention describes these antisense oligos as having cytostatic, dermatological and antiarteriosclerotic activities. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligonucleotide sequence is a chimeric phosphorothioate antisense oligo with 2' MOE wings and a decay gap, which is used to inhibit expression of mouse CTGF of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
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Sequence 20 BP; 11 A; 1 C; 0 G; 8 T; 0 U; 0 Other;

CTGF of the invention.

mouse

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Gaps ; 0

2; Indels

0; Mismatches

2824 ATATATATATATATATA 2842

17; Conservative

Matches

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Local Similarity

2 AAATATATATATATA 20

vivlemore401-10.rng

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Human NOVX polypeptide gene forward primer SEQ ID NO: 535.
   18-DEC-2003 (first entry)
                                                                                                                                                                                           WO2003000842-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-DEC-2001;
21-FEB-2002;
21-FEB-2002;
22-FEB-2002;
22-FEB-2002;
22-FEB-2002;
                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUN-2001;
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03-JUL-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JUN-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. rheumatoid arthritis). The present DNA sequence represents a 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to mouse TGF-beta receptor II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compound having a sequence targeted to a nucleic acid encoding
Transforming growth factor beta-receptor II, useful for preparing a
composition for treating hyperproliferative disorder e.g., lung, liver,
                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                  mouse; antisense oligonucleotide;
transforming growth factor beta receptor II; TGF-beta receptor II;
typerproliferative disorder; breast cancer; autoimmune disorder;
rheumatoid arthritis; 2.0-methoxyethyl gapmer;
phosphorothioate backbone; ss; murine.
                                                                                                                                                                                                                                                                         Mouse TGF-beta receptor II targeted antisense oligonucleotide #36
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0.4%; Score 15.8; DB 1; Length 20;
39.5%; Pred. No. 1.3e+03;
Ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
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                                                                2823 TATATACATATATAT 2841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1453 AAGGGTAACCTGCGGGAGT 1471
                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUN-2002; 2002WO-US019665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUN-2001; 2001US-00888361
                89.58;
                                                                                                                                                                           ADC65837 standard; DNA; 20
                                                                                                                                                                                                                                         (first entry)
                                17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        colon or gastric cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Murray SF, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-175279/17.
Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003000656-A2
                                                                                                                                                                                                                                         18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
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Matches
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20 AAGGCCAACCTGCAGGAGT 2

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ВР.

ADC10516 standard; DNA; 20

ADC10516 RESULT 1089
ADC10516/c
ID ADC10516
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AC ADC10516
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New isolated NOVX polypeptides and nucleic acid molecules useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory or CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Agee ML, Anderson DW, Berghs C, Casman SJ, Catterton E;
Dipippo VA, Edinger SR, Eisen A, Ellerman K, Gangolli EA;
Gerlach VL, Gorman L, Guo X, Herrmann UL, Hjalt T, Ji W, Kekuda F
Khramtsov NV, Li L, Liu X, Malyankar UM, Miller CE, Millet I;
Ort T, Padigaru M, Patturajan M; Pena CEA, Rastelli L, Rieger DK;
Rothenberg ME, Shenoy SG, Shimkets RA, Smithson G, Spaderna SK;
Spytek KA, Stone DJ, Vernet CAM, Zhong H, Zhong M, Alsobrook JP;
Burgess CE, Lepley DM;
                                                           neuroprotective, antiinflammatory; gene therapy; antisense therapy; thyromimetic, NOVX; pathology; cancer; diabetes; obssity; endocrine disorder; CNS disorder; inflammatory disorder; chromosome mapping; tissue typing; predictive medicine.
                                                cytostatic; antidiabetic; anorectic; cerebroprotective;
                                                                                                                                                                                                                                                                             2001US-0296575P.
2001US-0297414P.
2001US-0295573P.
2001US-0297567P.
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2001US-0298528P.
2001US-0299133P.
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2001US-0322297P.
2001US-0324669P.
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2001US-0299949P.
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2001US-0300883P.
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2001US-0341562P.
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2002US-0359122P.
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2002US-0359964P.
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ADH93257
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                                                The invention relates to novel isolated polypeptides, mature form of the polypeptide, a sequence that is 95% identical to the polypeptide or the polypeptide comprising one or more conservative substitutions. The NOVX polypeptide is useful for treating or preventing a pathology associated with the polypeptide e.g. disorders associated with aberrant expression or activity of the polypeptide, such as cancer, diabetes, obesity, and andocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing and predictive medicine. This sequence corresponds to a primer used to amplify and isolate the coding sequence for one of the polypeptides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compounds that hybridizes with nucleic acid molecules encoding farnesyl transferase beta subunit and inhibits the expression of farnesyl transferase beta subunit, useful for treating e.g. cancer or inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide; phosphorothioate linkage,
2'-O-methoxyethyl sugar moiety; 5-methylcytosine;
hyperproliferative disorder; cancer; ovarian carcinoma; adenocarcinoma;
colorectal cancer; pancreatic cancer; prostate cancer;
inflammatory condition; tumour formation; cytostatic; antiinflammatory;
antimicrobial; phosphorothioate oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human FT-beta subunit phosphorothioate oligonucleotide #46.
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
Example B; SEQ ID NO 535; 772pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1667 TGAAGATCGCAGACTTCGG 1685
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG93018 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                      the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAR-2004
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ID ADG9

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in cells or tissues and a method of treating an animal having a disease or condition associated with the FT-beta subunit. The compound is an antisense oligonuclectide, preferably a chimeric oligonuclectide, which comprises at least one modified internucleoside linkage which is a phosphorothioate linkage, at least one modified sugar moiety which is a 2'-O-methoxyethyl sugar moiety or at least one modified nucleobase which a 5-methylcytosine. The compound is useful in inhibiting the expression of the FT-beta subunit in cells or tissues. It can also be used for treating cells or conditions associated with the FT-beta subunit, such as typerprofilerative disorders, including cancer (such as ovarian carcinoma, adenocarcinoma, colorectal cancer, pancreatic cancer or prostate cancer) and inflammatory conditions. The antisense compounds can also be used as research agents, in diagnostics or for preventing or delaying infection, inflammation or tumour formation. This sequence represents a human farnesyl transferase beta subunit phosphorothioate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polynucleotide for detecting single nucleotide polymorphisms existing in human gene, contains isolated human gene having specified sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; gene sequence; single nucleotide polymorphism; SNP; disease diagnosis; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.8; DB 1;
Pred. No. 1.3e+03;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3727 AAACCGGCAGGTGCGATTT 3745
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 89.5%;
Matches 17; Conservative
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3778 AAGACACCTGGTTGCTAAC 3796

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The invention relates to a compound targeted to a nucleic acid molecule encoding the human farnesyl transferase beta (FT-beta) subunit and inhibits the expression of the (FT-beta) subunit, or specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding the FT-beta subunit. The invention also relates to a method of inhibiting the expression of the FT-beta subunit

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Human, antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiatethmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                         Sandrasagra A, Ka
L, Shahabuddin S;
                                                                                                 Human oligonucleotide sequence.
AAGGCACCTGGTTGCAAAC 1
                                                 BP
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                                                                                                                                                                                                                                         24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC
                                                 DNA; 20
                                                                                 (first entry)
                                                                                                                                                                                                                                                                         Li Y, Sar
Tang L,
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                                                 ABZ91732 standard;
                                                                                                                                                                                         WO200285308-A2.
                                                                                                                                                                         Homo sapiens.
                                                                                 17-OCT-2003
                                                                                                                                                                                                        31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                          ubiquinone
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                                                                  ABZ91732;
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                                                                                                                                                                                                                                                                          Nyce JW,
                                  RESULT 10
ABZ91732/
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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or Disclosure; SEQ ID NO 6974; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or deptenting levels of or entiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition.

Und inflammation, lung allergies, or a respiratory disease or condition. Specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 11 A; 0 C; 1 G; 8 T; 0 U; 0 Other;

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Gaps
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0
Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
 0.4%;
   Query Match 0.4
Best Local Similarity 89.5
Matches 17; Conservative
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19 rararrrrcarararar 1

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RESULT 1093 ABZ9173(

ABZ91730 standard; DNA; 20

ABZ91730;

17-0CT-2003

(first entry)

Human oligonucleotide sequence

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. antisense; lung dysfunction; nasal airway dysfunction;

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, I Miller S,

Aguilar D;

Pabalan J,

Katz E,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 6972; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a composition are spiratory, lung or malignant crespiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of or enclainflammatory steroid in a subject, for reducing levels of adenosine creeptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Once: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 8 A; 0 C; 2 G; 10 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 15.8; DB 1; Length 20; 19.5%; Pred. No. 1.3e+03; Indels 0; Mismatches 89.5%; Query Match
Best Local Similarity 89.55
Matches 17; Conservative

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3464 ATATATATCTATATATA 3482

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2823 TATATATACATATATAT 2841

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ABZ98620 standard; DNA; 20 BP.

ABZ98620;

(first entry) 17-0CT-2003

Human tryptase a oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Katz E, Pabalan J, Aguilar D; ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 13862; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an nasal airway dysfunction and a second active agent comprising an has antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is sufful for treating or preventing a respiratory, lung or maligiant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, the printed lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 5 A; 8 C; 5.G; 2 T; 0 U; 0 Other;

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Gaps
                                       ;
0
0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; ive 0; Mismatches 2; Indels
                                       17; Conservative
                      Best Local Similarity
        Query Match
                                       Matches
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1 CCCGCAGAAGTGCATCCA 19
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RESULT 1095

ΒÞ ABZ91729 standard; DNA; 20

ABZ91729,

ABZ91729;

(first entry) 17-0CT-2003

Human oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antisinflammatory steroid, ubiquinone, antisinflammatory, antiallergic, antisathmatic, hypotensive, immunosuppressive, cytostatic, gene therapy, antisense gene therapy, respiratory, lung, adenosine sensitivity, adenosine receptor, bronchodilation, bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar Katz E, Pabalan J, Li Y, Sandrasagra A, K Tang L, Shahabuddin S; Nyce JW, I Miller S,

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WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 6971; 872pp; English.

antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Socie: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an The invention relates to a novel pharmaceutical composition, which has a

Sequence 20 BP; 6 A; 1 C; 2 G; 11 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; tive 0; Mismatches 2; Indels Query Match Best Local Similarity 89.5% Warches 17, Conservative

2826 ATATACATATATATA 2844

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vivlemore401-10.rng

antipsoriatic; antiinflammatory; neuroprotective; ophthalmological; gastrointestinal, osteopathic; nephrotropic; gene therapy; multimeric molecular complex; transcrotric transport; paracellular transport; calcitonin; osteoporosis; renal failure; colitis; gastroentertils; inflammatory bowel disease; psoriasis; Alzheimer's disease; optic neuropathy; ophthalmoplegia; single chain antibody; sevSeR; linker; heavy chain region; ds.

Single chain antibody sPv5AF related linker #2.

(first entry)

20-MAY-2004

ADL99556;

ADL99556 standard; DNA; 20

RESULT 1097

ADL9955

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The invention relates to a method of selecting a single chain antibody (scFV) against a peptide target in a yeast by expressing a library of scFV tusion proteins in yeast cells, expressing a target fusion protein in the yeast cells expressing the scFV fusion proteins having either the DNA binding domain or the activation domain of the transcription activator which is not comprised in the scFV fusion proteins, and a target peptide, and selecting those yeast cells in which a reporter gene is expressed. Each scFV fusion protein comprises either an activation domain or a DNA binding domain of a transcription activator and a scFV having a heavy chain of a variable region (WH) of antibody whose sequence variable region (WH) of antibody whose sequence varies within the library independently of the VH antibody whose sequence varies within the library independently of the creporter gene is activated by a reconstituted transcriptional activator formed by binding of the scFV fusion protein to the target fusion are useful for protein. The methods and compositions of the present invention are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Selecting an scFv against a peptide target by expressing a target fusion protein having a DNA binding domain or activation domain of a transcription activator, useful for diagnosing, preventing and/or treating HIV infection and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for preventing and/or treating HIV infection and cancer. This sequence corresponds to tan oligonuclectide used to generate the scFv antibody of
                                                                                                                                                                                                                                                                     anti-HIV; cytostatic; virucide; single chain antibody; antibody; yeast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 0 C; 16 G; 0 T; 0 U; 4 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 4; 150pp; English.
                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-FEB-2002; 2002US-00071866.
08-FEB-2002; 2002US-00072031.
25-APR-2002; 2002US-00133978.
20 ATATACATACATATAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                              07-FEB-2003; 2003WO-US003763.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhu L;
                                                                                                               ADK17359 standard; DNA; 20
                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENE-) GENETASTIX CORP.
                                                                                                                                                                                                                                                                                             HIV; cancer; CCR5; BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pauling MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-731501/69
                                                                                                                                                                                                                                 Human CCR5 protein.
                                                                                                                                                                                                                                                                                                                                                                     WO2003066830-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the invention.
                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2003
                                                                                                                                                     ADK17359;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hua S,
                                                                                                               용
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Glynn JM;

Sheridan PL, Houston LL,

Hawley SB, Chapin S, WPI; 2003-898076/82.

06-SEP-2001; 2001US-00949039. 06-SEP-2001; 2001US-00949039.

US2003166160-A1

Synthetic.

04-SEP-2003,

(HAWL/) HAWLEY S B. (CHAP/) CHAPIN S. (SHEK/) SHERIDAN P L. (HOUS/) HOUSTON L L. (GLYN/) GLYNN J M.

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The invention describes a multimeric molecular complex comprising at least 2 compounds, each of which has at least one targeting element directed to a ligand that confers transcytotic or paracellular transporting properties to a molecular complex specifically bound to the ligand. Also described are: a compound comprising at least 2 targeting elements directed to the ligand; a protein compusion at least 2 targeting to composition composition to the ligand; a plant and an element directed to the ligand; a pharmaceutical composition comprising the compound; delivering a biologically active agent to an animal; transporting a biologically active agent to an animal; transporting a biologically active agent to an animal; treating a disease in an animal; or disease in an animal. The complex is useful for preparing a composition for diagnosing or treating diseases, e.g., osteoporosis, renal failure, colitis, gastroenteritis, inflammatory bowel disease, psoriasis, Alzheimer's disease, optic neuropathy or ophthalmoplegia. This sequence represents a linker associated with the isolation of heavy chain regions from the single chain antibody spv5AP polypeptide, that targets the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New multimeric molecular complex, useful for preparing a composition for diagnosing or treating e.g. osteoporosis, renal failure, colitis, gastroenteritis, inflammatory bowel disease, psoriasis or Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptide, that targets the liator of endocytosis, exocytosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and forward and reverse transcytosis in epithelial cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 0 C; 16 G; 0 T; 0 U; 4 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (plgR) mediator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Page 51; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2920 GGGCGGGCGTGGGGGGC 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polyimmunoglobulin receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 73.7
les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches
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Gaps

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Match 0.4%; Score 15.8; DB 1; Length 20; Local Similarity 73.7%; Pred. No. 1.3e+03; les 14; Conservative 4; Mismatches 1; Indels

Query Match

Best Loca Matches

2920 GGGCGGGCGTGGGGGGGC 2938

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The invention relates to an antisense oligonucleotide 8-30 nucleobases in length targeted to the 5'-untranslated region, translational start site, translational termination region or 3'-untranslated region of a nucleic acid molecule encoding a p38 mitogen activated protein kinase (MAPK), where the antisense compound inhibits the expression of the p38 MAPK. The antisense oligonucleotide is useful for inhibiting the expression of p38 MAPK in cells or tissues. It is also useful for treating an animal having a disease or condition associated with p38 MAPK, e.g. an inflammatory or an autolimnuc disease (e.g. rheumatoid arthritis) or a heart disease. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. The present sequence represents a p38 MAPK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides for modulating p38 mitogen activated protein kinase (MAPK) expression, useful for diagnosing, preventing or treating diseases associated with p38 MAPK, e.g. inflammation or heart
                                                                                                                                                                                    antisense; p38 mitogen activated protein kinase; p38 MAPK; inflammatory disease; autoimmune disease; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                   are 5-methyl cytosines"
                                                                                                                                                                                                                                                                                                                                                                            /mod_base= Other
/note= "2'-methoxyethoxy nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "2'-methoxyethoxy nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Phosphorothioate linkages"
                                                                                                                                                           Mouse p38 MAPK antisense oligonucleotide #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mckay R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; SEQ ID NO 65; 48pp; English.
                                                                                                                                                                                                                                                                                                                                  'note= "All cytosines
                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                    /mod_base= Other
                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= Other
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/mod_base= Other
 2 GGGSGGGSGGGSGGGS 20
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                                                                     ADM34276 standard; DNA; 20 BP
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15-AUG-2000; 2000US-00640101
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                                                                                                                              (first entry)
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                                                                                                                                                                                                                    heart disease; ss; mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16. .20
                                                                                                                                                                                                                                                                                           . 20
                                                                                                                                                                                                                                                                                                                                                                                                                         *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-898587/82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MONI/) MONIA B P. (GAAR/) GAARDE W A. (NERO/) NERO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MCKAY R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JS2003176383-A1
                                                                                                                                                                                                                                                                                        modified base
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                                                                                                                              03-JUN-2004
                                                                                                                                                                                                                                                Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monia BP,
                                                                                                 ADM34276;
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                                           RESULT 1098
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating or target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted toncleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pabalan J, Aguilar D;
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                                                                             Length 20
                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           Human Tryptase a-derived oligonucleotide SEQ ID 13862.
                                     Sequence 20 BP; 2 A; 10 C; 6 G; 2 T; 0 U; 0 Other;
                                                                           Score 15.8; DB 1;
Pred. No. 1.3e+03;
antisense oligonucleotide of the invention.
                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 15; SEQ ID NO 13862; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sandrasagra A, Ka
i, Shahabuddin S;
                                                                                                                                                             43 GGCCCCAGCGCTGCAGG 61
                                                                                                                                                                                  20 GTGCCGCAGCGGCTGCAGG 2
                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-APR-2002; 2002WO-US013143.
                                                                               0.4%;
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                                                                                                                                                                                                                                                                                                 ABD31651 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2004 (first entry)
                                                                         Query Match
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Li Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200285309-A2
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                                                                                                                                                                                                                                                                                                                                             ABD31651;
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                                                                                                                                                                                                                                                              RESULT 1099
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pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it.
   reduce the amount of target polypeptide present in the lungs. The
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
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0
      0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.38+03; Live 0; Mismatches 2; Indels
                                                                         1601 CCTCCCAGAAGTGCATCCA 1619
                                                                                                        cccccccacacaccarcca 19
                                         Conservative
Query Match
Best Local Similarity
Matches 17; Conserv
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BP ABD27959 standard; DNA; 20 (first entry) ABD27959; RESULT 1100

29-JUL-2004

respiratory tract inflammation; adenosine sensitivity; lung, cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; se; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; AA497002-derived oligonucleotide SEQ ID 6971.

Homo sapiens

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW,] Miller S,

WPI; 2003-093058/08.

to Pharmaceutical composition for treating asthma, has antisense objigonucleotide containing less percentage of adenosine, targeted tuncleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 6971; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The

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Composition of atarget polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The instruction or cancer and can be anti-sense to the corresponding mRNA. The instruction also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) adelivery of instructions for adding a carrier and for use of the kit. The composition of invention has antialergic, antialfammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to confine an antial and or availability, and/or bronchoconsfriction and/or lung confine and or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, charactery pain, cystic fibrosis, allergies thinitis, pulmonary contension, emphysema, chronic obstructive pulmonary disease, pulmonary content of the anti-sense olgose corresponding to thymidines present in the target RNA serves to prevent the breakdown of the unity beaution and the present in the target RNA serves to prevent the breakdown of the plant of the anti-sense of the plant of the partice manner.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g., lung, brain, heart, kidney, etc, prevent any unwanted effects due to it
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Sequence 20 BP; 6 A; 1 C; 2 G; 11 T; 0 U; 0 Other;

Gaps ; Score 15.8; DB 1; Length 20; Pred. No. 1.3e+03; 0; Mismatches 2; Indels Query Match 0.4%; Best Local Similarity 89.5%; Matches 17; Conservative '

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ABD27960 standard; DNA; 20 29-JUL-2004 (first entry) ABD27960; RESULT 1101

BP.

AA497002-derived oligonucleotide SEQ ID 6972.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Pabalan J, Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-093058/08.

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analgesic, hypotensive, immunosuppressive and cytostatic activity, is beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises olique and is administered to reduce the amount of target polypeptide present in the target mRNA or to reduce the amount of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction, respiratory distress syndrome, pain, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergiar bronchitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, t free adenosine into the system tissue environment and thereby, to Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and Claim 15; SEQ ID NO 6972; 763pp; English prevent any unwanted effects due to it lung, brain, heart, kidney, etc, bronchodilating agent.

Sequence 20 BP; 8 A; 0 C; 2 G; 10 T; 0 U; 0 Other;

0; Gaps Query Match 0.4%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.3e+03; Matches 17; Conservative 0; Mismatches 2; Indels

1 ATATGTATGTATATATA 19

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Homo sapiens

WO200285309-A2

31-OCT-2002

3464 ATATATCTATATATA 3482 ઠ В

ABD27962 standard; DNA; 20 BP 29-JUL-2004 (first entry) ABD27962; RESULT 1102 ABD27962 DXBXSXXXXXXXXXXXXXXXXXXXXX

AA497002-derived oligonucleotide SEQ ID 6974.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; bronchitis; pulmonary transplantation rejection; ss; primer.

Antisense; ss; calcineurin; regulatory subunit B; PPP3R1; immunosuppressive; nootropic; neuroprotective; protein phosphatase 3;

Antisense oligonucleotide targeting human PPP3R1 ISIS 206769

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This invention describes a novel composition (a) a first active agent, comprising oligomucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and control depletion or hyposecretion, when administered to a mammal. The oligomucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or reduce the amount of target polypeptide present in the lungs. The culticammation, allergies and/or bronchoconstriction and/or lung with a disease or condition such as pulmonary viseconstriction, cinflammation, allergies and/or bronchoconstriction, respiratory disease, pulmonary compagnate, pain, cystic fibrosis, allergic rhinitis, pulmonary content of the anti-sense oligos corresponding to the hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cransplantation rejection, pulmonary infections, branchines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenomine into the system e-g., lung, brain, heart, kidney, etc, tissue environment and thereby, to the prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                    invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                            oligonucleotide containing less percentage of adenosine, targeted unucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                          Pabalan J, Aguilar
                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating asthma, has antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.4%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.3e+03; Matches 17; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                          Katz E,
                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; SEQ ID NO 6974; 763pp; English.
                                                                                                                                                          Li Y, Sandrasagra A, K
Tang L, Shahabuddin S;
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                     23-APR-2002; 2002WO-US013143.
                                                                24-APR-2001; 2001US-0286036P.
                                                                                                             (EPIG-) EPIGENESIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                              bronchodilating agent.
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Miller S,
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The invention relates to a new compound comprising 8-50 nucleobases in
length targeted to a nucleic acid molecule encoding protein phosphatase 3
CPPPRIX, the regulatory subunit of calcineurin), where the compound
specifically hybridises with the nucleic acid and inhibits the expression
of PPPRIX, i.e. is an antisense oligonucleotide (AO). Also included are a
compound 8-50 nucleobases in length which specifically hybridises with at
least an 8-nucleobase portion of an active site on a nucleic acid
molecule encoding PPPRIX, a method of inhibiting the expression of PPPRIX
in cells or tissues (comprising contacting the cells or tissues with AO
contacting a disease or condition associated with PPPRIX (comprising
administering AO to the animal so that expression of PPPRIX is inhibited)
and a method of screening for an antisense compound (comprising
contacting a preferred target region of a nucleic acid molecule encoding
PPPRIX with one or more candidate antisense compounds which
an enclobase portion which is complementary to the preferred target
region, and selecting for one or more candidate antisense compounds which
inhibit the expression of a nucleic acid encoding PPPRIX is inhibited
and compositions of the present invention are useful for the diagnosis,
pervention and or activity of BPPRIX is not anti-
pervention and anticinal and antiched antisense diagnosis,
and selecting a preferred pervention are useful for the diagnosis,
and composition or activity of BPPRIX is not anti-
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and an active and a preferred and encoding and antiched antiched antiched and antiched and antiched and antiched and antiched antiched and antiched and antiched antiched and antiched and antiched and antiched and antiched antiched and antiched and antiched and antiched and antiched antiched and anticked with an antiched antiched and antiched and antiched and anticked and antiched and antiched and anticked and an
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                                                                                                                                                                                              /mod_base= OTHER
/note= "Phosphorothioate linkages and all cytidines are
-methylcytidines"
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  autoimmune disorder; aberrant calcium signaling; neurological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     like Alzheimer's disease. The present sequence is an anti-PPP3R1 antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                           /note= "2'-methoxyethyl nucleotides"
16. .20
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                                                                                                                        Location/Qualifiers
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                            OTHER
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                         Alzheimer's disease; human.
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*tag=
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modified_base
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Location/Qualifiers
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                                                                                                                                 Alzheimer's disease; human.
                             (first entry)
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modified base
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                                                                                                                                                              Homo sapiens.
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                             22-APR-2004
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ADI27315;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.8; DB 1;
89.5%; Pred. No. 1.3e+03;
tive 0; Mismatches 2;
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Best Local Similarity 89.59
....hes 17, Conservative
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The invention relates to a new compound comprising 8-50 nucleobases in length targeted to a nucleic acid molecule encoding protein phosphatase 3 (PPPSRI, the regulatory subunit of calcineurin), where the compound specifically hybridises with the nucleic acid and inhibits the expression of PPPSRI, i.e. is an antisense oligonucleotide (AO). Also included are a compound 8-50 nucleobases in length which specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding PPPSRI, a method of inhibiting the expression of PPPSRI is inhibiting the expression of PPPSRI is inhibitied) and an inhal having a disease or condition associated with PPPSRI (comprising administering AO to the animal so that expression of PPPSRI is inhibited) and method of screening for an antisense compound (comprising contacting a preferred target region of a nucleic acid molecule encoding PPPSRI with one or more candidate antisense compounds thaving at least an 8-nucleobase portion which is complementary to the preferred target
                                                                                                                                                                                                                                                                                                                                                                                                                    "mod_base= OTHER
'note= "Phosphorothioate linkages and all cytidines are 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of PPP3R1, such as autoimmune disorders, aberrant calcium signaling and Alzheimer's
                                                                                                                                                                                                                                           Antisense; ss; calcineurin; regulatory subunit B; PPP3R1; immunosuppressive; nootropic; neuroprotective; protein phosphatase 3; autoimmune disorder; aberrant calcium signaling; neurological disease;
                                                                                                                                                                                                            Antisense oligonucleotide targeting human PPP3R1 ISIS 206769.
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/mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
16. 20
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/mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
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region, and selecting for one or more candidate antisense compounds which inhibit the expression of a nucleic acid encoding ppp3R1). The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of ppp3R1, such as autoimmune disorders, conditions having aberrant calcium signaling and neurological diseases like Alzheimer's disease. The present sequence is an anti-ppp3R1 antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel single or multiple target oligonuclectide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
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                                                                                                                                                                                                                     Length 20;
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                                                                                                                                                                            Sequence 20 BP; 8 A; 1 C; 2 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                   Score 15.8; DB 1;
Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide associated to Tryptase-a #35.
                                                                                                                                                                                                                                                         0; Mismatches
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in S, Lu H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ60499 standard; DNA; 20
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                                                                                                                                                                                                           Query Match
Best Local Similarity 89.5'
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                   Therapy, insulin resistance, hyperglycaemia; type II diabetes mellitus;
human; antisense; integrin-linked kinase; ILK; p59ILK; ss.
(ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hyperglycemia, or type II to the mammal in need of the
                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                           Human integrin-linked kinase antisense oligonucleotide ISIS #109232
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/note= "2'-methoxyethyl (2'-MOE) nucleotides where
cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotides where
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                                                                              0.4%; Score 15.8; DB 1; Length 20;
39.5%; Pred. No. 1.3e+03;
                                                                                                        Indels
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/note= "Phosphorothioate backbone"
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/note= "2'-methoxyethyl (2'-MOE)
cytidines are 5-methylcytidines"
16..20
/*tag= c
                                                   Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating a mammal for insulin resistance, hype
diabetes mellitus comprises administering to t
treatment an integrin-linked kinase inhibitor.
                                                                                                       0; Mismatches
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                                                                                          89.58;
                                                                                                                                                                                                                              ADJ58861 standard; DNA; 20
                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                          Best Local Similarity 89.5
Matches 17; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                             invention.
                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                       ADJ58861;
                                                                               Query Match
                                                                                                                                                                                                    1106
                                                                                                                                                                                                    RESULT 1
ADJ58861
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Chimeric phosphorothioate oligonucleotide to target Navl.3 #4031.

(first entry)

20-MAY-2004

ADK76697;

ADK76697 standard; DNA; 20

RESULT 1108

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to meuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate but not limited to neonatal or infantile epilepsy; or ataxia. The sequence represents a chimeric phosphorothioate oligonucleotide with 2.7ME wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound targeted to a nucleic acid molecule encoding May1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                  Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                               Gaps
                                                                                                                                                                                                                                                                                                   Chimeric phosphorothioate oligonucleotide to target Navl.3 #3509.
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0
            Score 15.8; DB 1; Length 20; Pred. No. 1.3e+03;
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                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 15.8; DB 1;
89.5%; Pred. No. 1.3e+03;
0.4%; Sco...
89.5%; Pred. No. 1...
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                                                                             GAGGAGCTGGTGGAGGCTG 874
                                                                                                            GAGGAGCAGGTGGAGTG 20
                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2002; 2002US-0403416P.
                                                                                                                                                                                               ADK76175 standard; DNA; 20
                                                                                                                                                                                                                                                                 (first entry)
                            Local Similarity 89.5
les 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHAA ) PHARMACIA CORP
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                ADK76175;
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              Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              wew antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
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different regions of the human Nav1.3 RNA.
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Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                26-FEB-2004.
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                                                                                                                                                                                                                                                                                                                   Synthetic.
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Gaps

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2; Indels

0; Mismatches

1147 GAGCTGCCTGCCGACCCCA 1165

17; Conservative

Matches

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1 GAGCTGCATGCCGACCACA 19

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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1; inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immicrosomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuropropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; allabeles; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; acardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense compound, having a sequence targeted to a nucleic acid
                                                                                                                                                                                 /note = "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
chimeric; antisense oligonucleotide; phosphorothioate; human;
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                                                                                                                                                                                                                          /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                                                                                                                                                                                                                                 not\overline{e}= "2'-0-methoxyethyls"
                                                                                                                                         Location/Qualifiers
                                                                                                                                                                       /mod_base= OTHER
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/mod_base= OTHER
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                                                                                                                                                  modified base
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                                                                                                                                                                                                                                                modified base
                                                                                                          Homo sapiens
                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                              Gierse JK;
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of inhibits its expression; (2) a method of inhibiting the expression of mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (3) a method of inhibiting an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulators, and cardiavascular activities, and can cophthalmological, immunomodulators and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation Alzheimer's compound disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder. encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or Claim 4; SEQ ID NO 1011; 132pp; English. ischemia.

Sequence 20 BP; 8 A; 8 C; 4 G; 0 T; 0 U; 0 Other;

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0.4%; Score 15.8; DB 1; Length 20; 39.5%; Pred. No. 1.3e+03; Ved. No. 1.3e+03; Indels Ive. 0; Mismatches 2; Indels
                           89.58;
  Query Match 0.4
Best Local Similarity 89.5
Matches 17; Conservative
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Gaps

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; imPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antitinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9d34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
                                                                                                                                                                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        residues are 5-methylcytidines
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"

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    /*tag= a
    /mod_base= OTHER
    /note= "2'-0-methocyethyls"

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                                                                                                                   ВЪ.
2322 TGTGTGTGTGTGTGTGT
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                                                                                                                 ADM15524 standard; DNA; 20
                                                                                                                                                                                  01-JUL-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                  ADM15524;
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                                                                              RESULT 1110
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             antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, notropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric; antisense oligonucleotide; phosphorothioate; human; microscomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microscomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunoomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; vasotropic; ophthalmological; immunoomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
   having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1696.
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                                                                                                                                                                                                          Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                          Sequence 20 BP; 10 A; 8 C; 0 G; 2 T; 0 U; 0 Other;
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/note= "2'-O-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                         2315 GICTGTGTGTGTGTGTG 2333
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                                                                                                                                                                                                                                         Matches 17; Conservative
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                                                                                                                                                                                                                         Local Similarity
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ADM15509/
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WPI; 2004-305094/28.

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGBS-1). The human mpGBS-1 gene is located on chromosome 9, more specifically to 9444.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGBS-1, which specifically the britise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGBS-1 in cells or tissues; and (3) a method of treating an animal antisense condition associated with mpGBS-1. MpGBS-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidifammatory, neuroprotective, nootropic, antidifammatory introprotective, nootropic, antisinflammatory introprotective, nootropic, antisinflammatory introprotective, nootropic, antistinfismmatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; implibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer: disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                            encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                        New antisense compound, having a sequence targeted to a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 10 A; 8 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1. .5
/*tag= a
/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
/*tag= c
                                                                                                                           Claim 4; SEQ ID NO 1696; 132pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2315 GTCTGTGTGTGTGTGTG 2333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM14920 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 89.5
les 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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Homo sapiens.

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin B2 synthase (mPGBS-1). The human mPGBS-1 gene is located on chromosome 9, more specifically to operate of the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid mPGBS-1 and mPGBS-1, which specifically hybridise with the nucleic acid mPGBS-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGBS-1 in cells or tissues; and (3) a method of treating an animal companies of oligonucleotides and antisense oligonucleotides and antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamatory, neuroprotective, nootropic, antisense oligonucleotides and and ni gene therapy. The antisense compound to be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound can be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound confition associated with mPGBS-1 e.g., inflammation, Alzheimer's condition associated with mPGBS-1 e.g., inflammation, Alzheimer's diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; UCAM; tryptase a; tryptase b; DD24 A; DD24 B; DD24 C; DD24 B; DD24 C; DD24 B; DD24 C; DD24 B; DD26 B; DD24 C; DD25 C; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory disease; by disease; copp, allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match 0.4%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 8 A; 8 C; 3 G; 1 T; 0 U; 0 Other;
                     /note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 1107; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2326 TGTGTGTGCGTGTGTGT 2344
/mod base= OTHER
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                                                                                                                                                        25-SEP-2003; 2003WO-US030374.
                                                                                                                                                                                                  25-SEP-2002; 2002US-0413549P.
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                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                     WPI; 2004-305094/28.
                                                                 WO2004028458-A2.
                                                                                                             08-APR-2004
                                                                                                                                                                                                                                                                                             Gierse JK;
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Gaps ö

2; Indels

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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Edtaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(8) or expressed product(8), for the prevention and/or treatment of a respiratory or lung disease. The cligonucleotides are useful for reducing or inhibiting expression of a configuration or treating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor for asthma and/or lung allergies associated with impaction or an inflammation, allergy, anthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary disease (CDPD), cystic fibrosis (CP), chronic obstructive pulmonary disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                Aguilar D, Miller S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 1.3e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; SEQ ID NO 1355; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                  A, Tang L,
Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1601 CCTCCCAGAAGTGCATCCA 1619
                                                                                                                        25-JUL-2003; 2003US-00627930.
                                                                                                                                                            23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                                                    SANDRASAGRA A.
                                                                                                                                                                                                                                                                                                            SHAHABUDDIN S
                                                                                                                                                                                                                                                                                                                                                                                                   Shahabuddin S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-293804/27.
                                                                                                                                                                                                                                                     TANG L.
AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                   NYCE J W
                                                                                                                                                                                                                                                                                         MILLER S
                                                   US2004049022-A1.
                                                                                                                                                                                                                                                                                                                             LU H.
CONG H.
                                                                                      11-MAR-2004
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                                                                                                                                                                                                                 NYCE/)
                                                                                                                                                                                                                                                                                                                                                CONG/)
                                                                                                                                                                                                                                                                                                            SHAH/)
                                                                                                                                                                                                                                      SAND/)
                                                                                                                                                                                                                                                                                           MILL/
                                                                                                                                                                                                                                                                                                                           LUHHI/
                                                                                                                                                                                                                                                         TANG/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   asthma
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Matches
8.46
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ADO51745 standard; DNA; 20 BP.

ADOS1745 ID ADOS XX

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ADAMIS; metagirdin; MDC1S; a disintegrin and metalloproteinase domain 15; diagnosis; inflammation; therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention is directed to antisense oligonucleotides targeted to ADAMIS (otherwise known as metagirdin, MDCIS, and a disintegrin and metalloproteinase domain 15) and which modulate the expression of ADAMIS. The invention is useful for diagnosing and treating diseases associated human the expression of ADAMIS such as inflammation. The present sequence is human ADAMIS target to oligonucleotide. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                            New compound targeted to a nucleic acid molecule encoding ADAM15 and inhibits the expression of ADAM15, useful for modulating the expression of ADAM15 or for diagnosing or treating, e.g. inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= Other
/note= "Phosphorothioate backbone. All cytidines are 5-
methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 9 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; Antisense Therapy; Gene Therapy; nidogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human nidogen antisense oligonucleotide #52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 53; 38pp; English
                                    Human ADAM15 target oligonucleotide #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chediak-Higashi syndrome; ss; probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cocation/Qualifiers
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/*tag= a
/mod_base= Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  196 GCTGAGGACACAGGTGTGG 214
                                                                                                                                                                                                                                                                                                                                                        Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                          21-NOV-2002; 2002US-00302028
                                                                                                                                                                                                                                                                             21-NOV-2002; 2002US-00302028
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12-AUG-2004 (first entry)
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Matches 17; Conservative
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/*tag=
                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                        Dean NM,
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-399722/37.
                                                                                                                                                                 US2004102392-A1
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modified base
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                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                    Bennett CF,
                                                                                                                                                                                                      27-MAY-2004
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                                                                                                ADAM15; metagirdin; MDC15; a disintegrin and metalloproteinase domain 15; diagnosis; inflammation; therapy; human; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                /note= "Phosphorothioate backbone where all cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention is directed to antisense oligonucleotides targeted to ADAMIS (otherwise known as metagirdin, MDC15, and a disintegrin and metalloproteinase domain 15) and which modulate the expression of ADAMIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compound targeted to a nucleic acid molecule encoding ADAM15 and inhibits the expression of ADAM15, useful for modulating the expression of ADAM15 or for diagnosing or treating, e.g. inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention is useful for diagnosing and treating diseases associated with expression of ADAMIS such as inflammation. The present sequence is human ADAMIS antisense oligonucleotide. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                        (2' -MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "2' -methoxyethyl (2' -MOE) nucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                       Human ADAM15 antisense oligonucleotide ISIS #173627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 4 C; 9 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHBR
/note= "2' -methoxyethyl
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                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                 5-methyl cytidines"
                                                                                                                                                                                                                                                                                             /mod_base= OTHER
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/mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dobie KW;
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ADO51782 standard; DNA; 20
                                  (first entry)
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004102392-A1
                                                                                                                                                                                                                                                        modified base
                                                                                                                                                                                                                                                                                                                                                    modified base
                                                                                                                                                                                                                                                                                                                                                                                                                            modified base
                                                                                                                                                                                   Homo sapiens
                                    12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAY-2004
                                                                                                                                                                                                      Synthetic.
ADO51745;
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Gaps

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/note= "2'-methoxyethyl (2'-MOE) nucleotides"

AD051782

ADOS1782/ ID ADOS XX AC ADOS XX

RESULT 1115

g

Best Loca Matches

FFFFFFFF X B X B X B

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The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an xenotransplant rejection or mechanical organ replacement rejection, in an individual. The method is also useful in assessing the immune status of individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or thugal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                       Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a compound targeted to a nucleic acid molecule encoding the human ABCCS polypeptide. The compound is an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ABCC5; 88; antisense oligonucleotide; phosphorothioate linkage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotide compound that inhibits expression of ABCC5, usefu
for preparing a composition for treating hyperproliferative disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 3 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2'-O-methoxyethyl sugar moiety; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human ABCC5 DNA antisense oligonucleotide #44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                 Claim 58; SEQ ID NO 755; 1762pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; SEQ ID NO 54; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2320 TGTGTGTGTGTGTGTGT 2338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 rérérérecriérécérer 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-DEC-2002; 2002US-00319893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADP44428 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-SEP-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
  WPI; 2004-400724/37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g., cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-2004
                                                                                                                                                the genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADP44428;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADP44428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to antisense oligonucleotides which are targeted to, and inhibit the expression of, a nucleic acid molecule encoding nidogen. The antisense oligonucleotides are useful for treating a disease or condition associated with nidogen, such as Chediak-Higashi syndrome. They are also useful in research and diagnostics for modulating the expression of nidogen. The present sequence represents a human nidogen antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds, particularly oligonucleotides targeted to a nucleic acid encoding nidogen, useful for treating diseases associated with nidogen, e.g. Chediak-Higashi syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transplant rejection; immune system; rheumatoid arthritis; lupus;
inflammatory bowel disease; multiple sclerosis; HIV; AIDS; 88; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                     /note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prentice J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 8 A; 2 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set 1 left PCR primer for marker probe #91.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ly N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 15; SEQ ID NO 64; 91pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2822 GTATATATACATATATA 2840
16. .20
/*tag= c
/mod_base= Other
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                                                                                                                                                                                                                                                                                                                 19-NOV-2002; 2002US-00300611
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                                                                                                                                                                                                                                                         19-NOV-2002; 2002US-00300611
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20-DEC-2002; 2002US-00325899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP10746 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17; Conservative
                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fry K,
                                                                                                                                                                                                                                                                                                                                                                                                                              Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-389192/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004042346-A2.
                                                                                                                                          US2004097451-A1
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Rosenberg S;
  modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-AUG-2004
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                                                                                                                                                                                                 20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                 Chiang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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Matches
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IID ADP1

XX ADP1

XX ADP1

XX IIC-A

XX IIC-A
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Gaps

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oligonucleotide that specifically hybridises with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothiotete linkage, at least one modified sugar moiety, preferably a 2'-C-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of the human ABCC5 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence repersents an antisense oligonucleotide targeted to DNA encoding the human ABCC5 polypeptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                              .
0
                                                                                                                                                                                                                                                                                                                 Score 15.8; DB 1; Length 20;
Pred. No. 1.3e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                         0.4%;
                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 89.5%,
Thes 17; Conservative
       88888888888888888
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ADP44503 standard; DNA; 20 BP ADP44503; RESULT 1119 ADP44503

2687 AGGCTTTCCCACTTCCCAC 2705

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2 AGGITITICCACTÍCCCAC 20

Human ABCC5 DNA antisense oligonucleotide target region #41. 09-SEP-2004 (first entry)

Human; ABCC5; ss; antisense oligonucleotide; phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; hyperproliferative disorder; cancer; cytostatic.

Homo sapiens

US2004115649-A1.

17-JUN-2004.

12-DEC-2002; 2002US-00319893

12-DEC-2002; 2002US-00319893.

(ISIS-) ISIS PHARM INC

Dobie KW;

WPI; 2004-449386/42.

useful New oligonucleotide compound that inhibits expression of ABCC5, usefu for preparing a composition for treating hyperproliferative disorder, e.g., cancer.

Example 15; SEQ ID NO 129; 57pp; English.

The invention relates to a compound targeted to a nucleic acid molecule encoding the human ABCC5 polypeptide. The compound is an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage i.e. a phosphorothicate linkage, at least one modified sugar moiety, preferably a 2'-0-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methyloycosine. The antisense compounds are useful for modulating the expression of the human ABCC5 polypeptide and in preparation of a composition for treating hyperproliferative disorders, e.g. cancer. This sequence reperseents a human ABCC5 DNA antisense oligonucleotide target region of the invention.

Seguence 20 BP; 7 A; 3 C; 7 G; 3 T; 0 U; 0 Other;

RESULT 1121

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                                                                                                                                                                                                                                                                                                                                                 Human; 88; antisense; STAT 6;
signal transducer and activator of transcription; transcription factor;
rheumatoid arthritis; obesity; allergy; autoimmune disorder;
chromosome 12q13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide compound that inhibits expression of STAT 6, useful for preparing a composition for treating e.g. autoimmune disorders.
                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
                                       .;
0
0.4%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.3e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.4%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.3e+03; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for the antisense oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 8 A; 8 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                               Human STAT 6 antisense target region #22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 108; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dobie KW
                                                                           2687 AGGCTTTCCCACTTCCCAC 2705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2317 CTGTGTGTGTGTGCG 2335
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                                                                                                         19 AGGTTTTTCCACTTCCCAC
                                                                                                                                                                                                                                                                          (first entry)
                    Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shanahan WR, Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                            09-SEP-2004
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                                                                                                                                                                                                                                      ADP68501;
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     Query Match
                                                                                                                                                                   RESULT 1120
ADP68501/c
                                                                                                          셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "Phosphorothioate backbone and all cytidines are 5
-methylcytidines"
                                                                                                                                                                                                                                                          Human; 88; antisense; STAT 6; signal transcription; transcription factor; rheumatoid arthritis; obesity; allergy; autoimmune disorder; chromosome 12q13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide compound that inhibits expression of STAT 6, useful for preparing a composition for treating e.g. autoimmune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.8; DB 1; Length 20; Pred. No. 1.38+03; 0; Mismatches 2; Indels
                                                                                                                                                                                                                Human STAT 6 antisense oligonucleotide ISIS153790.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 3 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          not\bar{e}= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 15; SEQ ID NO 40; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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/*tag= b
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                               ADP68433 standard; DNA; 20 BP
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/*tag= a
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Best Local Similarity 89.5'
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                                                                                           ADP68433;
ADP6 8433
ADP6 8433
ADP6 ADP6
ACA ADP6
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deoxyribonucleic acid, major groove, ethanoamino group;
aziridinylcytosine, cross-linking group, tumour necrosis factor; ss.
                                                                                                                                                                                                                                                                                                   /*tag= d
/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
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/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
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/note= "N-methyl-8-oxo-2'-deoxyadenine"
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/note== "N-methyl-8-oxo-2'-deoxyadenine'
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/note= "N-methyl-8-oxo-2'-deoxyadenine'
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/note= "N-methyl-8-oxo-2'-deoxyadenine'
                                                                                                          Cross-linking oligomer 218 for targetting human TNF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= k
/mod_base= OTHER
/note= "N4N4-ethanocytosine"
                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                          *tag= c
mod_base= OTHER
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/mod_base= OTHER
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91US-00640654.
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                                                      AAQ20036 standard; DNA; 21
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14-JAN-1991;
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                                    RESULT 1122
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New sequence-specific non-photo-activated crosslinking agents - bind to the major groove of duplex DNA and are esp. useful for treating latent infections e.g. HIV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.4%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 1.4e+03; Matches 17; Conservative 0; Mismatches 2; Indels
/*tag= h
/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine'
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/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
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/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine'
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/mod_base= OTHER
/note= "N4N4-ethanocytosine"
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14-JAN-1991;
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                                                                                                                                          New sequence-specific non-photo-activated crosslinking agents - bind to the major groove of duplex DNA and are esp. useful for treating latent infections e.g. HIV.
                                                                                                                                                                                                                                                               The sequence is designed to target the Human tumour necrosis factor beginning at nucleotide 1137 and to covalently cross-link to it via the N4N4-ethanocytosine group. See also AAQ20031-Q20038
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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aziridinylcytosine; cross-linking group; tumour necrosis factor; ss.
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/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
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mod_base= OTHER
note= "N-methyl-8-oxo-2'-deoxyadenine'
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/note== "N-methyl-8-oxo-2'-deoxyadenine'
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/note= "N-methyl-8-oxo-2'-deoxyadenine
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                                                                                                                                                                                                                                                                                                                                                Seguence 21 BP; 10 A; 1 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "N4N4-ethanocytosine"
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                                                                                                                                                                                                                         Example 4; Page 25; 42pp; English.
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/mod_base= OTHER
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_base= OTHER
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/mod_base= OTHER
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                                                           Matteucci MD, Krawczyk S;
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                   (GILE-) GILEAD SCIE INC
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3468 ATATCTATATATATTT 3486
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08-APR-1991;
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17-APR-1991;
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                                                                                                                                                                                                                                                                                                                                            This PCR primer was used together with AAQ26545 to amplify a 150bp stretch of DNA from the RING11 gene containing a polymorphism. Sequencing of the enrire RING11 sequence from EDNA clones from B 1 ymphoblastoid cell lines revealed a silent single base pair mutation, and 3 single base pair mutation, and 3 single base pair mutarion, and 3 single base pair mutaristed region. 2 of the coding sequence substitututions lead to amino acid substitutions, one of which changed the putative stop codon, and lengthened the protein by 17 amino acids. Caussoid controls were also analysed by oligonucleotide typing with probes AAQ26546-51
                                                                                                                                                                                                                       and diagnosis of immune disorders and screening of new immunosuppressants and immuno:enhancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligomer TNF218 for forming triplex with HUMTNFAA target duplex.
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/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 1.48+03; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 3 A; 4 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                       Powis SH;
                                                                                                                (IMCR ) IMPERIAL CANCER RES TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                       Kelly AP, Glynne R,
                                                                                                                                                                                                                                                                                                          Example 2; Page 40; 101pp; English.
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/mod_base= OTHER
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 GGATCGCACAGTGCTGGTG 21
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                  91WO-GB002278.
                                                                          91GB-00019711
                                                        90GB-00027520
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(first entry)
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                                                                                                                                                                                         WPI; 1992-250030/30.
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modified_base
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                  19-DEC-1991;
                                                                                                                                                       Trowsdale J,
                                                        19-DEC-1990;
                                                                          16-SEP-1991;
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07-DEC-1992
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/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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/mod_base= OTHBR
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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0.4%; Score 15.8; DB 1; Length 21;
Best Local Similarity 89.5%; Pred. No. 1.4e+03;
Matches 17; Conservative 0; Mismatches 2; Indels
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/mod_base= OTHER
/note= "OTHER= N4 N4 ethanocytosine"
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                                                                                                                                                                                                                        /*tag= g
/mod_base= OTHER
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91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
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Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV;
malignancy; hepatitis; inflammation; 8s.
                                                                                      Oligomer TNF219 for forming triplex with HUMTNFAA target duplex.
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/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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|mod_base= OTHER
|note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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|mod_base= OTHER
'note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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mod_base= OTHER
'note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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                                                                                                                                                          Location/Qualifiers
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/mod_base= OTHER
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/mod_base= OTHER
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/mod_base= OTHER
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/mod_base= OTHER
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                   AAQ30388 standard; DNA; 21
                                                          (revised)
(first entry)
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                                                        25-MAR-2003
07-DEC-1992
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                                                                                                                                      Synthetic
                                      AAQ30388;
RESULT 1126
           AAQ30388/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer; polymerase chain reaction; PCR; amplification; heavy chain; immunoglobulin; Ig; variable; framework; region; production; recombinant; antibody; B cell; diagnosis; therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                           New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis, herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer VH3 for heavy chain variable region framework region 1 cDNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 15.8; DB 1; Length 21;
89.5%; Pred. No. 1.4e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                   Matteucci MD, Milligan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 9 A; 2 C; 0 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                      Claim 12; Page 70; 77pp; English.
91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546..
91US-00686547.
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Matches 17; Conservative
                                                                                                                                  Froehler B, Krawczyk S,
                                                                                                      (GILE-) GILEAD SCI INC
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 18-JAN-1991;
08-APR-1991;
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                               17-APR-1991;
                                           17-APR-1991;
                                                                        27-SEP-1991;
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                                                          7-APR-1991
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Welschof M;

Terness P,

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encoding an immunoglobulin (Ig), heavy chain, variable region, framework region 1, which corresponds to residues 1-7 of the Ig sequence. The primer can be used in a novel method for the production of recombinant antibodies, comprising the selection of B cells from a lymphocyte fraction, isolation of mRNA from individual B cells, reverse transcription of the mRNA into cDNA, amplification of the cDNA by PCR and cloning and expression of the cDNA. The recombinant antibodies can be used for diagnosis and/or therapy, while the method avoids the need for intermediate separation of light and heavy chains and gene library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a PCR primer for IgG used in the present invention which describes a human monoclonal antibody (hMAb). The CDNA encoding the hMAb can be used for commercial production of the hMAb. The CDNA was isolated from an antitetanus toxin human monoclonal antibody producing hybridoma cell strain TTG6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New cDNA encoding human monoclonal antibody - useful for production of antibody by hybridoma techniques commercially.
                                                                                                                                                       The present sequence is a primer for the PCR amplification of the cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, monoclonal antibody; hybridoma cell strain TTG6; hWAb; IgG; antitetanus toxin; immunoglobulin G; PCR primer; ss.
                                                                   Prodn. of recombinant antibodies - by amplification, cloning and expression of cDNA generated from B-cell mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.4%; Score 15.8; DB 1; Length 21; Best Local Similarity 81.0%; Pred. No. 1.4e+03; Matches 17; Conservative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunoglobulin G PCR primer VH3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          853 GAGGAGGAGCTGGTGGAGGCT 873
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                                                                                                                      Claim 5; Page 8; 11pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-138233/13.
                                WPI; 1997-088250/09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thiazide sensitive cotransporter and ATP sensitive potassium channel genes - useful for developing products for the diagnosis and treatment of ion transport disorders, e.g. Gitelman's Syndrome or Bartter's Syndrome.
                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                     ion transport; Gitelman's syndrome; Bartter's syndrome; hypokalaemic alkalosis; hypocalciuria; hypomagnesemia; diagnosis; therapy; SSCP; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primers hTSCex1A forward and reverse (see AAV40565 and AAV40566,
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19.5%; Pred. No. 1.4e+03;
ve 0; Mismatches 2; Indels
                              3; Indels
 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gitelman's syndrome and other ion transport disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 2 A; 11 C; 4 G; 4 T; 0 U; 0 Other;
Score 15.8; DB 1;
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                             Human TSC gene exon 1 forward primer hTSCex1A
                              1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 51; 105pp; English.
                                                               853 GAGGAGCTGGTGGAGGCT 873
                                                                                           21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3404 GTTTCCAGGGAGGGCCGG 3422
                                                                                               1 GAGGTGCAGCTGKTGGAGTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 GTGTCCAGGGAGGGGCCAG
 0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-US023553.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96US-00778052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89.58;
                                                                                                                                                                             AAV40565 standard; DNA; 21
                                                                                                                                                                                                                                              (first entry)
 Query Match
Best Local Similarity 81.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lifton RP, Simon DB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-388029/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYYA ) UNIV YALE
                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9829431-A1
                                                                                                                                                                                                                                              21-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUL-1998.
                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                               AAV40565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Gaps

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RESULT 1130

Sequence 21 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 1 Other;

ACC58780/

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New estrogen-receptor gene derived from bluegill animal encoding estrogen-receptor, useful for evaluating estrogen-receptor activity regulation ability of subject.
                                                                                                                                                                                                                                                                                                                The invention relates to a novel gene encoding an oestrogen-receptor protein. The invention may be useful for measuring the oestrogen-receptor activity regulation ability of a subject and is efficient in evaluating the oestrogen-receptor activity regulation ability of a chemical substance in an environment. The current sequence is that of the bluegill sunfish oestrogen-receptor-related PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention comprises isolated human gene sequences and PCR primer sequences which can be used to detect single nucleotide polymorphisms (SNPs). The DNA sequences of the invention are useful for detecting SNPs existing in human genes and for the diagnosis of human disease. The present DNA sequence represents a human gene PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polynucleotide for detecting single nucleotide polymorphisms existing in human gene, contains isolated human gene having specified sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.8; DB 1; Length 21; 89.5%; Pred. No. 1.4e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human, gene sequence; single nucleotide polymorphism; SNP; disease diagnosis; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 3 A; 11 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 1906; 529pp; Japanese.
                                                                                                                                                                                                                                                                           Example 2; SEQ ID NO 24; 30pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             646 GTGGAGGTGAATGGCAGCA 664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 gresagercaaccagra 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADH94069 standard; DNA; 21 BP
                                       11-JAN-2002; 2002JP-00004395.
                                                                            11-JAN-2002; 2002JP-00004395.
                                                                                                                 (SUMO ) SUMITOMO CHEM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-DEC-2001; 2001JP-00377637.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-DEC-2001; 2001JP-00377637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human gene PCR primer #914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                          WPI; 2003-819594/77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-819215/77.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-APR-2004
  15-JUL-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADH94069;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is that of an antisense oligonucleotide that is targeted to an accessible region of P2X3 mRNA. P2X3 is a purinoreceptor that is implicated in pain sensitivity and bladder volume reflexes. The antisense oligonucleotide specifically hybridises within an accessible region of the P2X3 mRNA in its native state (see also ACC58776 and ACC58777), thereby reducing P2X3 production. Cells or tissues are contacted with the antisense oligonucleotide, resulting in an inhibition of pain sensory neurons, or an increase in bladder capacity. The method can be used to alleviate pain, e.g. in patients suffering from chronic pain, and to treat disorders of urine storage and voiding, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotide for inhibiting the production of the purinoreceptor P2X3 and for treating pain or disorders of urine storage and voiding, such as overactive bladder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oestrogen receptor; regulation; environment; bluegill sunfish; ss; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                          Purinoreceptor; receptor; P2X3; analgesic; uropathic; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.8; DB 1; Length.21;
Pred. No. 1.4e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hart LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer SEQ ID 24 used to amplify bluegill sunfish DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 21 BP; 2 A; 3 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhang H,
                                                                                                                                     Purinoreceptor P2X3 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stone LS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 17; Page 10; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                716 CTAACACCACCGACAAGGA 734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTAACCTCACCGACAAGGA 3
780/c
ACC58780 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                    08-NOV-2002; 2002WO-US036000
                                                                                                                                                                                                                                                                                                                                                                           09-NOV-2001; 2001US-0337338P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arvidsson UNG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF92493 standard; DNA; 21
                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              (NEUR-) NEUROMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-430663/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        overactive bladder
                                                                                                                                                                                                                                                        WO2003040339-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shuster SJ,
                                                                                              26-AUG-2003
                                                                                                                                                                                                                                                                                           15-MAY-2003
                                                                                                                                                                                                                 Synthetic
                                                     ACC58780;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF92493;
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NAME OF THE PROPERTY OF THE PR

Sequence 21 BP; 1 A; 2 C; 9 G; 9 T; 0 U; 0 Other;

Lepomis macrochirus

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JP2003199580-A.

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Gaps

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The invention relates to a novel artificial animal cell expressing a variant androgen receptor. The variant androgen receptor is obtained by introducing a reporter gene, base sequence for initiating transcription, to the chromosome of the animal cell, where amino acid sequence of the variant androgen receptor is identical to that of a normal androgen receptor and has different amino acids at positions 646, 781 or 895 of a fully defined sequence of 918 amino acids as given in the specification. The artificial animal cell is useful for evaluating a substance capable of modulating the activity of the variant androgen receptor, which involves contacting a test substance with the animal cell, measuring the index value which has correlation and the expression value of the reporter gene or its quantity, and evaluating the ability of substance to modulate the variant androgen receptor based on the measured index value and the expression value of the reporter gene or its quantity. The method is useful for retrieving the substance which has the ability of modulating the variant androgen receptor. The variant androgen receptor regulator. The is useful for exercise in a substance that binds with the variant androgen receptor regulator. This
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polynucleotide sequence represents a probe used in the exemplification of
                                        Gaps
                                                                                                                                                                                                                                                                                                                                 Human androgen receptor related southern hybridisation probe, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                       variant androgen receptor; reporter gene; chromosome; animal cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel artificial animal cell expressing variant androgen receptor obtained by introducing reporter gene, sequence for initiating transcription to cell, useful for evaluating substance modulating
                                        ;
0
  Length 21;
                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 9 C; 5 G; 2 T; 0 U; 0 Other;
Score 15.8; DB 1;
Pred. No. 1.4e+03;
                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 8; 61pp; Japanese.
                                                                          2318 TGTGTGTGTGTGTGCGT 2336
                                                                                                                 1 rererereserererer 19
                                                                                                                                                                                                                 BP.
  0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-JUN-2002; 2002JP-00168252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUN-2002; 2002JP-00168252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SUMO ) SUMITOMO CHEM CO LID
                                                                                                                                                                                                                                                                                                                                                                                             regulator; human; ss; probe.
                                                                                                                                                                                                               ADL96726 standard; DNA; 21
                                                                                                                                                                                                                                                                                          20-MAY-2004 (first entry)
                                        17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-113321/12.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2004008142-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-2004.
                                                                                                                                                                                                                                                      ADL96726;
                                                                                                                                                                           RESULT 1133
                                        Matches
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Human; NOVX, PCR, ss; metabolic disorder; diabetes; obesity; infectious disease; anorexia; cancer; neurodegenerative disorder; hataimer's disease; Parkinson's disease; immune disorder; haematopoietic disorder; antidiabetic; anorectic; antimicrobial; anabolic; eating disorder; cytostatic; neuroprotective; nootropic; antiparkinsonlan; antianaemic; primer.
21 GCTGGTGGTGCTGGAAGCC 3
                                                 ВЪ
                                                                                                                                                                                                                                                                                       2001US-0322816P.
2001US-0323519P.
2001US-0323631P.
                                                                                                                                                                                                                       06-SEP-2002; 2002US-00236392.
                                                                                                                                                                                                                                                                        2001US-0318765P.
                                                                                                                                                                                                                                                                               2001US-0322781P.
                                                                                                                                                                                                                                                                                                                2001US-0323636P.
                                                                                                                                                                                                                                                                                                                      001US-0324969P
                                                                                                                                                                                                                                                                                                                                      2001US-0324990P.
                                                                                                                                                                                                                                                                                                                                               2002US-0357303P.
                                                                                                                                                                                                                                                                                                                                                       2002US-0360973P
                                                                                                                                                                                                                                                                                                                                                               2002US-0366131P.
                                                                                                                                                                                                                                                                                                                                                                       2002US-0367753P.
                                                                                                                                                                                                                                                                                                                                                                                002US-0369479P.
                                                                                                                                                                                                                                                                                                                                                                                       2002US-0379532P
                                                                                                                                                                                                                                                                                                                                                                                                002US-0381664P
                                                 ADN96330 standard; DNA; 21
                                                                                (first entry)
                                                                                               Human NOVX PCR primer #93.
                                                                                                                                                                                                                                                                                                                               2001US-032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MALYANKAR U M.
BURGESS C E.
GERLACH V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MILLER C E.
SHENOY S G.
PATTURAJAN M.
PENA C E A.
TCHERNEV V T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANDERSON D W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CATTERTON E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PADIGARU M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TAUPIER R J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GUSEV V Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        LI L.
GORMAN L.
SPYTEK K /
KEKUDA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                               ZHONG M.
                                                                                                                                                                                        US2004067490-A1.
                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                        20-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUN-2002;
                                                                                01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                20-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                02-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                .7-MAY-2002;
                                                                                                                                                                                                        08-APR-2004
                                                                ADN96330;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MALY/)
(BURG/)
(GERL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PATT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ZHON/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       LILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TAUP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PADI/)
GUSE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TCHE/
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Gaps

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0.4%; Score 15.8; DB 1; Length 21; 19.5%; Pred. No. 1.4e+03; ve 0; Mismatches 2; Indels

89.28;

Local Similarity 89.5 tes 17; Conservative

Query Match Best Local Si Matches 17;

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Zhong M, Li L, Gorman L, Jericon B, Miller LE, Anderson DW, Vernet CAM, Catterton B, Miller LE, Batturajan M, Pena CEA, Tchernev VT, Padigaru M, Gusev VY; Malyankar UM, Burgess CE, Gerlach V, Casman SJ, Rieger DK; Grosse WM, Smithson G, Peyman JA, Starling G, Rothenberg ME; Larochelle WJ, Shimkets RA, Crabtree J, Rastelli L, Voss EZ; Larochelle WJ, Shimkets RA, Millet I, Macdougall JR, Ellerman K;
                                                                                                                                                                                                                                                                                                  Example C; SEQ ID NO 393; 552pp; English.
                      SMITHSON G.
PEYMAN J A.
STARLING G.
ROTHENBERG M E.
ILAROCHELLE W J.
SHIMKETS R A.
                                                                                                                               MACDOUGALL J R
                                                                                   RASTELLI L.
VOSS E Z.
BOLDOG F L.
EDINGER S R.
MILLET I.
                                                                   SHIMKETS R /
                                                                                                                                                                                                                                              WPI; 2004-355290/33.
                                                                                                                                                 CHAPOVAL A.
                                                                                                                                          ELLERMAN K
       (RIEG/)
(GROS/)
(SMIT/)
(STAK/)
(STAK/)
(LARO/)
(SHIM/)
(CRAB/)
(CRAB/)
(CRAB/)
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(CRAB/)
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(CRAB/)
(CRAB/)
                                                                                                                                        (ELLE/)
(CHAP/)
                                                                                                                                (MACD/
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New isolated polypeptide, useful for treating or preventing a pathology associated with the polypeptide, e.g. diabetes, infectious disease, cancer, neurodegenerative disorders or Alzheimer's disease.

isolated nucleic acids can be used to express the novel proteins, to detect novel mRNA or a genetic lesion in a novel gene and to modulate its activity. It can also be used in gene therapy for treating or preventing a pathology associated with the protein or nucleic acid. The disorders include metabolic disorders, diabetes, obesity, infectious diseases, anorexia, cancer, neurodegenerative disorders, Alzheimer's disease, Parkinson's disease, immune disorders and haematopoietic disorders. This sequence represents a PCR primer used in analysis of expression of a human NOVX polynucleotide of the invention. invention relates to human NOVX polypeptides and polynucleotides. The

Sequence 21 BP; 1 A; 7 C; 5 G; 8 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 15.8; DB 1; Length 21; 19.5%; Pred. No. 1.4e+03; ve 0; Mismatches 2; Indels 89.58; 17; Conservative Best Local Similarity Query Match Matches

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3554 TAGCCTGGACTGCTACCTT 3572

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ADJ93185 standard; DNA; 22 BP.

ADJ93185;

Human G-coupled receptor protein HGPRBMY30 gene probe.

(first entry)

06-MAY-2004

ds; gene; immunosuppressive; cardiant; antiinflammatory; cytostatic; anti-HIV; antirheumatic; antiarthritic; antiibacterial; antiseborrheic; dermatological; antipsoriatic; neuroprotective; noctropic; antiparkineonian; antidiabecic; ophthalmological; antiasthmatic; antidepressant; neuroleptic; hypotensive; tranquilizer; hypertensive;

ABL56893 standard; DNA; 30 BP.

ABL56893 ID ABL5

RESULT 1136

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The invention relates to an isolated human G-protein coupled receptor, HGPRBMY30 polypeptide or a sequence having 95% identity to the above mentioned sequences. (I) is useful for preventing or treating a medical condition, selected from an immune disorder; a cardiovascular disorder; an inflammatory disorder in which G-protein coupled receptors are either disorder; a neutral disorder; a med secondar; a metabolic disorder; a neutral disorder; an endocrine disorder; cesticular cancer; a neutral disorder; an endocrine disorder; cesticular cancer; an entangly sorders, and consider; as neutral disorder; (I) and (II) are also useful for detecting, prognosing, preventing, treating, and/or ameliorating the diseases such as hematopoietic and pulmonary disorders, Alzheimer's, Parkinson's diseases, diabetes, dwarfism, color blindness, retinal pigmentosa, as thema, expression, schizophrenia, sleeplessness, hypertension, anxiety, stress, renal failure, acute heart failure, hypotension, obesity, and extivity. (I) and (II) are useful for modulating signal transduction crivity. (I) and (II) are useful for modilating signal transduction activity. (I) and (II) are useful for modilating signal transduction crivity. (I) and (II) are useful for modilating signal transduction activity. (I) and (II) are useful for modilating signal transduction, size and shape, to change a mammal's mental state by physical state by influencing biorhythms, caricadic rhythms, copy planet, weight, hair color, eye color, skin, percentage of adipose tissue, pigmentation, size and shape, to change a mammal's mental state capabilities, hormonal or endocrine levels, appetite, librio, memory, erress, or or brist cognitive qualities. This sequence corresponds to a recent for pain the part of the rogalities. This sequence corresponds to a recent for pain the sequence corresponds to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human G-protein coupled receptor, HGPRBMY30 polypeptide useful for preventing and treating e.g. immune disorders, cardiovascular disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Pred. No. 1.4e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 5 A; 12 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; SEQ ID NO 88; 343pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probe for the novel HGPRBMY30 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Feder JN, Mintier GA, Ramanathan C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2596 CCCTCCCACACCCAAAGCT 2614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CCCTGCCACACCACAGCT 19
                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2001; 2001US-0294411P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%;
                                                                                                                                                                                                                                                                                                                                           30-MAY-2002; 2002WO-US017085.
                                                                                                                                                                                                                                                                                                                                                                                                                               (BRIM ) BRISTOL-MYERS SQUIBB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17; Conservative
                                                                                                                                                              myocardial infarction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-140445/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                     WO200296946-A1
                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                               05-DEC-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches
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ABL56893;

Synthetic.

Kurane Koyama

vivlemore401-10.rng

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Measurement of nucleic acids, using a nucleic acid probe and analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Probe; polymorphism detection; mutation detection; disease diagnosis; microbial identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.8; DB 1; Length 30; 74.1%; Pred. No. 1.9e+03;
                                                                                     ss; fluorochrome; nucleic acid probe; fluorescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30 BP; 4 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                             (KANK-) KANKYO ENG KK.
(KEIZ-) KEIZAI SANGYOSHO SANGYO GIJUTSU SOGO KEN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3259 AGATATTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe poly f for assaying nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 ATATATTTTTTTCTTTTTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 17; 34pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL95890 standard; DNA; 30 BP.
                                                                                                                                                                                                                                                                                                       20-APR-1999; 99JP-00111601.
24-AUG-1999; 99JP-00236666.
30-AUG-1999; 99JP-00242693.
01-FEB-2000; 2000JP-00028896.
                                                                                                                                                                                                                                                                                                                                                                                                                        BIOI-) BIOINDUSTRY KYOKAI SH
                                                                                                                                                                                                                                                                20-APR-2000; 2000JP-00120097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUN-2001; 2001WO-IB001147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-SEP-2000; 2000JP-00292483
11-APR-2002 (first entry)
                                         Poly f nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.4
Best Local Similarity 74.1
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-134193/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the obtained data
                                                                                                                                                                          JP2001286300-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200208414-A1
                                                                                                                                 Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUN-2000;
03-AUG-2000;
                                                                                                                                                                                                                      16-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                           (KANK-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to the determination of the concentration of a nucleic acid target, using a fluorescently labeled probe which produces reduced fluorescence measuring a fluorescently labeled probe which produces The method comprises measuring the reduction in emission caused by hybridisation. The new method is particularly used to quantify target mucleic acids by a real-time polymerase chain reaction, e.g. for quantifying microbial cells in co-cultures or symblotic systems, for detecting gene mutations or polymorphisms, and for analysing melting curves of target nucleic acids to determine a Tm value. Methods of the invention allow target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and no materials are introduced that inhibit amplification by Taq polymerase (so
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            conventional PCR conditions can be used). The specificity of PCR is kept high (amplification of primer dimers is delayed), and the limit of quantitation is reduced. Complex propers are not needed, and amplification can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation coefficient than conventional graphs so more accurate quantitation is possible. The current sequence represents a synthetic deoxyribooligonucleotide that was used for investigating the base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining the concentration of a target nucleic acid, useful e.g. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yokomaku T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                            Concentration; quantification; mutation detection; polymorphic; polymerase chain reaction; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.8; DB 1; Length 30;
Pred. No. 1.9e+03;
0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kamagata Y, Kurata S, Yamada K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 30 BP; 4 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                     Synthetic deoxyribooligonucleotide poly f.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GICCTITIT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NIBI-) JAPAN BIOINDUSTRY ASSOC.
(AGEN ) AGENCY OF IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                selectivity of a target nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Page 21; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA97617 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                            99JP-00111601.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%;
ilarity 74.1%;
Conservative C
                                                                                                                                                                                                                                                                                                                                   20-APR-2000; 2000EP-00108643
                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kanagawa T,
Furusho K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3259 AGATATTTATT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                         26-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                               20-APR-1999;
                                                                                                                                                                                                                                           EP1046717-A2
                                                                                                                                                                                                                                                                                       25-OCT-2000.
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Query Match

Best Loc Matches

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ABA97617

ABA97617 ID ABA9 XX AC ABA9 XX

RESULT 1137

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Gaps

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Indels

0; Mismatches

nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid

SKGGGG

Sequence 31 BP; 4 A; 1 C; 0 G; 26 T; 0 U; 0 Other

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Gaps

.. 0

3; DB 1; Length 31; 2e+03; 7; Indels

Score 15.8; D Pred. No. 2e+0 0; Mismatches

Query Match
Best Local Similarity 74.1%;
Matches 20; Conservative

and

124/c AAD27124 standard; RNA; 37

RESULT 1140 AAD27124,

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The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleotide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
                            Yamada K;
                                                                                           Fluorescently-labeled nucleic acid probes for assaying nucleic acids their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
                            Torimura M, Kurata S,
                                                                                                                                                                                                                                                                                                                                                                       Score 15.8; DB 1; Length 30; Pred. No. 1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                  7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; fluorochrome; nucleic acid probe; fluorescence.
                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 4 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                             3259 AGATATTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATATATTTTTTTTTTTTTTTTTTTT 29
                                                                                                                                                              Example 12; Page 60; 152pp; Japanese.
                           Kanagawa T, Kamagata Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                     ch 0.4%;
1 Similarity 74.1%;
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA97621 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Poly j nucleotide sequence
(KANK-) KANKYO ENG CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-APR-2002 (first entry)
                                                                   WPI; 2002-195876/25
                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP2001286300-A.
                                                                                                                                                                                                                                                                                                                   invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                           Kurane R,
Yokomaku T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       m
                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABA97621;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1139
                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                  the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA9762
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RNA template, (AU)3 used to direct RNA synthesis by HCV RNA polymerase.

(first entry)

09-APR-2002

AAD27124;

Hepatitis C virus, HCV replicase, non-structural protein 5B, NS5B, lead compound; RNA polymerase; ss.

Unidentified US6322966-B1

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99US-00309670. 99US-00309670.

27-NOV-2001.

11-MAY-1999; 11-MAY-1999;

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The present invention relates to an assay system for hepatitis C virus

(HCV) replicase activity. The assay system comprises an RNA template that
has an unstable, small stemloop at the 3' end capable of forming a copy-
back structure, a HCV non-structural protein 5B (NSSB), ATP, GTP, CTP,

and UTP nucleoside triphosphates (NTPS), where one of the NTP is
and UTP nucleoside triphosphates (NTPS), where one of the NTP is
and UTP nucleoside triphosphates to the identification of optimal
NSSB. The invention also relates to the identification of optimal
properties of an RNA template for copy-back self-priming RNA synthesis of
HCV. This activity can be used to screen for anti-HCV replicase compounds
to to characterise the biological relevance of lead compounds. The
CY NSSB polymerase mechanistically and kinetically and for designing
CY NSSB polymerase mechanistically and kinetically and for designing
System of the invention is useful for detecting HCV replicase activity.
The nucleic acid synthesised by NSSB is detected by evaluating an
utoradiograph of reaction products separated by gel electrophoresis. The
present sequence is RNA template, (AU)3 used to direct RNA synthesis by
CNA polymerase proteins of HCV, BVV and polivirus. This sequence is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Fig 2A; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhong W, Hong Z, Lau JYN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-096587/13.
                                                                                                                                                                                                                                                                                                                                                                                                                           (HONG/) HONG Z.
(LAUJ/) LAU J Y N.
                                                                                                                                                                                                                                                                                                                                                                                                       (ZHON/) ZHONG W.
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Measurement of nucleic acids, using a nucleic acid probe and analysis

(KANK-) KANKYO ENG KK. (KEIZ-) KEIZAI SANGYOSHO SANGYO GIJUTSU SOGO KEN

WPI; 2002-134193/18

BIOI-) BIOINDUSTRY KYOKAI SH

99JP-00236666. 99JP-00242693.

99JP-00111601

20-APR-1999;

24-AUG-1999; 99JP-0023666. 30-AUG-1999; 99JP-00242693. 01-FEB-2000; 2000JP-00028896.

20-APR-2000; 2000JP-00120097.

16-OCT-2001

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This invention relates to a method for measuring nucleic acids using

Example 5; Page 17; 34pp; Japanese

the obtained data.

Query Match

X S

Best Loca Matches

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The present sequence is one of a number of oligonucleotides which were used for the generation of RNA-protein fusions, including fusions having a myc epitope tag. The RNA-protein fusions comprise a protein covalently linked to the 3' end of its own mRNA. This is accomplished by synthesis and in vitro or in situ translation of an mRNA molecule with a peptide acceptor attached to its 3' end. The RNA-protein fusions are incubated under high salt conditions to produce a protein library. This method is useful for improving or altering axisting proteins, as well as for isolating new proteins and nucleic acid or small molecule targets. It may also be used to improve human or humanised single-chain antibodies for the treatment of a number of diseases. The method is useful for the isolation of proteins with specific binding properties, for screening complaining and claims ends on the basis of protein-protein interactions. Unlike prior art, the new method does not rely on manntaining the integrity of an mRNA-ribosome:nascent chain ternary complex, which is very fragile and is therefore of limited use. The method does not rely on topological links between the protein and the nucleic acid so that the information of the protein is retained and can be recovered in readable, nucleic acid form
                  fusion; protein library; protein isolation; DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Producing protein or DNA libraries which are useful for improving existing proteins, by in vitro translating protein coding sequences to produce RNA-protein fusions and incubating these protein fusions under
                                                                                                                                                                                                                                   /*tag= b
/mod_base= OTHER
/note= "attached to puromycin, a peptide acceptor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

0.4%; Score 15.8; DB 1; Length 42;
Best Local Similarity 74.1%; Pred. No. 2.3e+03;
Matches 20; Conservative 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 42 BP; 31 A; 4 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3262 TATTTTATTTGCTTTGTCCTTTTTCAG 3288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           37 rriririririririririririririricas 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 44; 121pp; English.
                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ä
                                                                                                                                                                                                                                                                                                                                                                                                                  01-FEB-2000; 2000WO-US002589
                                                                                                                                                                       /*tag= a
/label= RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV48091/c
ID AAV48091 standard; DNA; 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Szostak JW, Roberts RW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            high salt conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-533022/48.
                    RNA-protein fusio
gene cloning; ss.
                                                                                                                                                                                                                                                                                                                               WO200047775-A1.
                                                                                                                                                                                                                     modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                          17-AUG-2000,
                                                                                    Synthetic
                                                                                                                                Key
misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV48091;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1143
셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Transcription factor from Aspergillus oryzae which regulates alpha-amylase promoter - useful for producing heterologous proteins, especially medicinal proteins or enzymes, in filamentous fungi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequence is that of PCR primer oligodT which was used in the analysis of a transcription factor (amyR) which regulates the expression of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    alpha-amylase; promoter; filamentous fungi; transcription factor; expression; control; production; heterologous polypeptide; medicinal; industrial enzyme; PCR primer; amyR; ss.
                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                             Aspergillus oryzae alpha-amylase transcription factor PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                              Match 0.4%; Score 15.8; DB 1; Length 37; Local Similarity 74.1%; Pred. No. 2.2e+03; es 20; Conservative 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.8; DB 1; Length 4
65.7%; Pred. No. 2.3e+03;
cive 0; Mismatches 12; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 41 BP; 2 A; 1 C; 2 G; 36 T; 0 U; 0 Other;
                    Sequence 37 BP; 32 A; 0 C; 2 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3300 TTCTATAGGATTTTTTTAGGAGATTTATTTTTT 3334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rrrigiaagcrirrrrrrrrrrrrrrrrrrrrrrrrr 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA-protein fusion oligonucleotide 43-P[CUG].
                                                                                                                                                     3259 AGATATTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                              Example 1; Page 25; 64pp; English
                                                                                                                                                                                                                                                                                                          AAV03013 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97WO-DK000305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96DK-00000740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA94319 Standard; DNA; 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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Best Local Similarity 65.77
"...hes 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NOVO ) NOVO-NORDISK AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          alpha-amylase promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-100998/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
Aspergillus oryzae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Christensen T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9801470-A1.
                                                                                                                                                                                                                                                                                                                                                                                                 17-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA94319;
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RESULT 1142

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Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor segment. The present sequence is an example of a SSR primer, which is flanked amplified fragment encledtides. The method represents a modified amplified fragment encledtides. The method represents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and
                                                                                                                                                                                                                                                                                                                                                                                     Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
  amplified fragment length polymorphism assay; microsatellite region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                     genetic trait marking; germplasm comparisons; 5'-anchored; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of a polymorphism site in a target gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.6; DB 1; Length 18; 83.3%; Pred. No. 1.2e+03; ive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gene inhibitor, cancer, allele specific inhibitor;
proliferative disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 7 A; 8 C; 0 G; 0 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 76; 173pp; English
                                                                                                                                                                                                                                                                  M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2337 GTGTGTGTGTGTGCAC 2354
                                                                                                                                                                                                                                                                  ප
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18 GTGTGTGTGTGTGTGVHV 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF80086 standard; DNA; 21 BP.
                                                                                                                                                                                  95WO-US015150.
                                                                                                                                                                                                                          94US-00346456,
                                                                                                                                                                                                                                                                  (DUPO ) DU PONT DE NEMOURS &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-00045054.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 83.3
nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (VARI-) VARIAGENICS INC.
                                                                                                                                                                                                                                                                                                        Vogel JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   germplasm comparisons
                                                                                                                                                                                                                                                                                                                                           WPI; 1996-277795/28.
                                                                                                  WO9617082-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                  21-NOV-1995;
                                                                                                                                                                                                                          28-NOV-1994;
                                                                                                                                                                                                                                                                                                          Morgante M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6200754-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUN-2001
                                                                                                                                            06-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF80086;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The Oligonucleotides AAV48087, AAV48089-V48091 and AAV48096-V48098 and the selection of a specific protein or RNA, by in vitro or in situ translation of candidate RNA-protein fusions. These were used in translation of specific proteins to produce RNA-protein fusions, then selecting specific RNA protein fusions. The method is used to select proteins (or DNA encoding them) having altered properties, e.g. for identification of new binding reagents, to identify improved human antibodies or new enzymes. These proteins are potentially useful in diagnosis and therapy, or as industrial catalysts. The methods allow many rounds of selection and amplification to be performed, resulting in enrichment of even very rare molecules and allowing isolation of proteins that bind specifically to almost any compound or catalyse almost any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Selection of specific protein by screening protein-RNA fusions generated in vitro or in situ - useful for, e.g. identifying enzymes and antibodies with altered properties, potentially useful as catalysts or for therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                           In situ translation, RNA-protein fusion, binding reagent, antibody, industrial catalyst, ss; rRNA/DNA hybrid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.8; DB 1; Length 43;
Pred. No. 2.4e+03;
0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5'-anchored simple sequence repeat primer BDB(CA)7.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 43 BP; 31 A; 4 C; 6 G; 0 T; 1 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3262 TATTITATTIGCTTIGICCTTTTCAG 3288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              37 rrrrrrrrrrrrrrrrrrrrrrrrrrrrr
                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 39; 94pp; English.
                                                                                                                                                                                                                              /*tag= a
/note= "Puromycin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Liu R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT27913 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                98WO-US000807
                                                                                                                                                                                                                                                                                                                                                                                                        97US-0035963P
97US-0064491P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ch 0.4%;
1 Similarity 74.1%;
20; Conservative
                                     Oligonucleotide 43-P [CUG]
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JAN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Szostak JW, Roberts RW,
                                                                                                                                                                                                                          *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-414032/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis
                                                                                                                                                                                                                                                                                                                                                                  14-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                        21-JAN-1997;
06-NOV-1997;
27-0CT-1998
                                                                                                                                                                                                                                                                                   WO9831700-A1
                                                                                                                                                                                                                                                                                                                           23-JUL-1998
                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                              Identifying inhibitor active on conditionally essential gene (EG) subject to loss of heterozygosity in cancer, useful in cancer treatment, involves determining two alleles of EG and testing potential allele specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cancer; ss; single nucleotide polymorphism; human; SNP; CEG; LOH; conditionally essential gene; alternative allele; loss of heterozygosity; antiproliferative treatment; DNA excision repair protein; ERCC5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/standard_name= "Single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 15.6; DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 8 A; 2 C; 5 G; 5 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence surrounding polymorphism target 1528.26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 1.4e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymorphism sites of various target genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stanton VP;
   Stanton VP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                               Disclosure; Fig 2; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1294 GTGAAGATGCTGAAAG 1309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS58215 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-00045054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 GTGAASATGCTGAAAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-FEB-2001; 2001US-00782837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (VARI-) VARIAGENICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Housman DE, Ledley FD,
   Ledley FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-066906/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chromosome 13q33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JS2002127714-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
   Housman DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2002.
                                                                                                                                                                                                                                  inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS58215;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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ABS58215
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create one but less than all of the alternative alleles. Also included care: (1) a method of identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on CEG, and where the content of cancer, where the inhibitor is active on CEG, and where the cancer involves and where the petient is suffering a potential allele determining at least two alleles of the gene, testing potential allele determining at least two alleles of CEG, where the patient is suffering from cancer, involves: (a) identifying a patient heterozygous for the gene, or the determining whether cancer cells in the patient have undergone loss concer, involves: (a) identifying a patient heterozygous for the gene, or the determining whether cancer cells in the patient have undergone loss of first allelic form of CEG (but not a second), where the portion comprises a sequence variance site; and (4) selecting an patient for a first allelic form of CEG (but nor a patient wifering a patient for comprises a sequence variance site; and (4) selecting an anipoliferative treatment with an antipoliferative treatment, or selecting an comprises a sequence variance site; and (4) selecting an anipoliferative treatment colls in a potential patient concerc calls of the patient have only the second allelic form, of the gene; is more active than a second allelic form, and determining where calls of the patient have only the second allelic form, the gene, where if the somatic cells are heterozygous and the cancer calls in a patient is suitable for treatment with the antiproliferative treatment is suitable for the patient of that the patient is suffable for treatment with the antiproliferative treatment is suitable for CEG, for inhibiting growth of a cell (involves abelient of the patient is suitable for CEG, for inhibiting processing and containing and concert where the patient with the subjecting at least one inhibitor active on allele of CEG, in allele of CEG, in allele of CEG, in the method inhibit separates between allelic form of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
Inhibitor for treating cancer, is active on allelic form of conditionally essential gene which has two alternative alleles in a population, and targets alternative alleles.
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                                                                                                                                                The invention relates to an inhibitor which is active on an allelic form
                                                                                                                                                            of a conditionally essential gene (CEG) comprising at least two alternative alleles in a population, and where the inhibitor targets at least one but less than all of the alternative alleles. Also included are: (1) a method of identifying an inhibitor notentially useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polymorphism in an allele of the target gene, DNA excision repair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.6; DB 1; Length 21; 93.8%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 8 A; 2 C; 5 G; 5 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
                                                                                                Disclosure; Fig 2; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein, ERCCS (chromosome 13q3)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1294 GTGAAGATGCTGAAAG 1309
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(first entry)
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Best Local Similarity
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05-JUN-1995
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                                                                                                                                                                                                                                                             PCR primers (AAQ72427-30) used for the mutagenesis of the tumour necrosis factor-alpha (TNF-a) (AAQ73431) to produce the mutein TNFa(D143N-A145R) (AAR62479). The resultant fragments were ligated into the expression plasmid pDS56/RBSII The expression of the wild type or mutein proteins is regulated by the lac repressor present on the plasmid pREP4. The gene encoding the protein is mutated at specific sites resulting in series of mutated proteins (AAR65493.103). The biological activities of TNF are mediated via specific receptor of mol. wt. 55 and 75 kDa called p55-TNF-R and p75-TNF-R respectively. The mutated proteins presented have a higher affinity for the human p75-TNF receptor than for the p55-TNF receptor. The mutated proteins can be used in a variety of therapeutic or diagnostic applications including cancer therapy, treatment of obseity, septic shock or bacterial meningitis. (Updated on
                                                                                                                                                                                                            New human TNF-a muteins with higher affinity for p75-TNFR - useful e.g. for cancer therapy, treatment of obesity and toxic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
receptor; affinity; therapeutic; diagnostic; cancer therapy; cancer; obesity; septic shock; meningitis; PCR; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Acridinium ester; probe; target; rabbit globin; screening assay; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4; Indels
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                                                                                                                                                                    Stueber D;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 3 A; 4 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.6; DB 1;
31.8%; Pred. No. 1.5e+03;
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                                                                                                                                                                  Lesslauer W, Loetscher H,
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                                                                                                                                                                                                                                             Example 1; Page 35; 53pp; English.
                                                                                                                                           (HOFF ) HOFFMANN LA ROCHE & CO AG
                                                                                                                                                                                                                                                                                                                                                                                                 treatment of obesity, septic shoc 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ85327 standard; DNA; 22 BP.
                                                                                                  94EP-00104154.
                                                                                                                        93EP-00810224.
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Best Local Similarity 81.8%;
Matches 18; Conservative
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                                                                                                                                                                                        WPI; 1994-311810/39
                                                                                                                       29-MAR-1993;
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21-AUG-1995
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                                                      EP619372-A1
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                                 Synthetic
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                                                                                                                                                              Oligo:nucleotide screening assay method - to determine the ability of the oligo:nucleotide to form a hybrid with a target nucleic acid sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 7 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1803 CGTCTGGTCCTTTGGGGTCCTG 1824
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                                                                                                                                                                                                                                                                                          Example; Page 24; 42pp; English.
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                                                                                 WPI; 1995-075251/10.
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Best Local Similarity
Matches 18; Conserv
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Nelson NC,
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Query Match 0.4%;
Best Local Similarity 81.8%;
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                 (neurofibromatosis type 2) gene from various human tumours. NP2 is an autosomal, dominantly inherited disorder characterised by multiple tumours of the central nervous system, predominantly bilateral vestibular cumonsomal acoustic neuromas) of the sighth cranial nervo. Other schwannomas and presentle lens opacities. The NP2 gene, mapped to chromosomal region 22412 between the loci D2281 and D2228, acts a tumour suppressor. The NP2 gene is alternatively spliced resulting in three different isoforms encoding three different proteins, merlin isoforms I.

III, which are likely to have differing functions. Merlin stands for moesin-ezrin-radixin like protein, so called due to substantial homology with these three proteins. The NP2 gene isoforms and proteins encoded by the under suppressors and can be used to suppress tumour growth, as can the cDNA sequence in gene therapy applications. Antibodies raised against merlin proteins are useful as tumour targetting agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a PCR primer for DNA encoding the protein of the invention, which induces signal transduction of the vascular endothelial cell growth receptor KDR (kinase insert domain-containing receptor) by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KDR signal transduction inducer protein; human; diabetic retinopathy; vascular endothelial cell growth receptor; abnormal neovascularisation; kinase insert domain-containing receptor; solid tumour proliferation; gene therapy; metastasis; chronic rheumatoid arthritis; psoriasis; retinopathy; retinopathy of prematurity; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer for KDR signal transduction inducer protein coding sequence.
                                      are PCR primers used for the isolation of the NF2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KDR signal transduction inducing protein and antibodies to it.
                                                                                                                                                                                                                                                                                             0.4%; Score 15.6; DB 1; Length 22; 81.8%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                       4; Indels
                                                                                                                                                                                                                                                                     Seguence 22 BP; 4 A; 10 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                               GCCGAGGAGGTGGTGGAGG 871
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                                                                                                                                                                                                                                                                                                                                                               GCTGAAGAGGAGCTGGTTCAGG 1
              Disclosure; Col 14; 46pp; English
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                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX84681 standard; DNA;
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                                                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-DEC-1998;
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Matches
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binding to its intracellular domain. The protein can be used in the investigation, diagnosis and treatment (including gene therapy) of diseases in which abnormal neovascularisation takes place, such as solid tumour proliferation, metastasis, chronic rheumatoid arthritis, psoriasis and retinopathy (including diabetic retinopathy and retinopathy of prematurity). It may be used as a screen for candidate KDR signal transduction inhibitors for therapeutic use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KDR signal transduction inducer protein; human; diabetic retinopathy; vascular endothelial cell growth receptor; abrormal necvascularisation; kinase insert domain-containing receptor; solid tumour proliferation; gene therapy; metastasis; chronic rheumatoid arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer for KDR signal transduction inducer protein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KDR signal transduction inducing protein and antibodies to it.
                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 15.6; DB.1; Length 22; 11.8%; Pred. No. 1.5e+03; ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    retinopathy; retinopathy of prematurity; PCR primer; ss
                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 7 A; 2 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 22 BP; 6 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           orematurity). It may be used as a screen ror
transduction inhibitors for therapeutic use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1288 GTAGCCGTGAAGATGCTGAAAG 1309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; Page 65; 82pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTAGCAGTCATGATGTTGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-JP005612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97JP-00343474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KYOW ) KYOWA HAKKO KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.4*;
Best Local Similarity 81.8*;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX84680/c
ID AAX84680 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shibuya M, Yabana N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-405033/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHIB/) SHIBUYA M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9931238-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX84680;
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Gaps

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Score 15.6; DB 1; Length 22; Pred. No. 1.5e+03; 0; Mismatches 4; Indels

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PCR primer; ss.
                                       WO200147959-A2
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                     Homo sapiens.
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                                                           05-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-SEP-2001
                                                                                                                                                                      Jager D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA168502;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1154
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes neurofibromatosis type 2 (NF2) transcript isoforms. NF2 polynucleotides can be used for diagnosing NF2 diseases, for inhibiting growth of tumnours associated with NF2 mutations (including expression from cDNA introduced in gene therapy vectors) and to raise antibodies (useful as tumour targeting agents, since specific isoforms are often tumour-specific) and as immunoassay reagents for detecting NF2-expression products. NF2 is a tumour suppressor protein, and so has muticancer activity. The present sequence represents a PCR primer for mouse NF2, from an example of the present invention
                                                                                                                                                                                                                                                                                                                                                                  Proteins from neurofibromatosis type 2 transcript isoforms - used for diagnosis or inhibition of tumors, and generation of antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cancer associated antigen; ING1; tumour suppressor; cancer; vaccine;
                                                                                                                                                             Human; neurofibromatosis type 2; NF2; tumour suppressor; cancer; diagnosis; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for cDNA encoding a human cancer associated antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.6; DB 1; Length 22;
31.8%; Pred. No. 1.5e+03;
ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 4 A; 10 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                         Mouse neurofibromatosis type 2 PCR primer 3ml
         1288 GTAGCCGTGAAGATGCTGAAAG 1309
                                                                                                                                                                                                                                                                                                                           Seizinger BR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               850 GCCGAGGAGGAGCTGGTGGAGG 871
                      22 gergaagagagergerreage 1
                                                                                                                                                                                                                                                                                                      (BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                                                                                                                                                                                                              Example; Col 14; 45pp; English.
                                                                               AAX04299 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH28492 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                96US-00628145
                                                                                                                                                                                                                                                                                   94US-00179738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81.8%;
                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18; Conservative
                                                                                                                                                                                                                                                                                                                          Bianchi AB, Kley NA,
                                                                                                                                                                                                                                                                                                                                              WPI; 1999-166715/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 18; Conserv
                                                                                                                      16-APR-1999
                                                                                                                                                                                                                                                                04-APR-1996;
                                                                                                                                                                                                                                                                                    .0-JAN-1994;
                                                                                                                                                                                                                        US5872214-A.
                                                                                                                                                                                                                                           16-FEB-1999
                                                                                                                                                                                          Synthetic.
                            22
                                                                                                   AAX04299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                      Mus sp.
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AAH28492
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PCR primers AAH28492-93 were used to amplify cDNA encoding a human cancer associated antigen, which is a variant of the ING1 gene. The ING1 gene is a tumour suppressor gene candidate. The cancer associated antigen polynucleotides and polypeptides are useful for screening for the possible presence of a pathological condition in a subject such as cancer. The cancer associated antigen polypeptides are useful for
                                                                                                                                                                                                                                                                         Chen Y;
                                                                                                                                                                                                                                                                                                                                                                                         Isolated cancer associated nucleic acid molecule identified by SEREX (serological identification of antigens by recombinant expression cloning) technique, useful in nucleic acid based therapies to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps

    L. monocytogenes iap gene cluster III PCR primer iap-1047-III-R.

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0
                                                                                                                                                                                                                                                                         Gure A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; iap gene; p60 protein; detection; infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 15.6; DB 1; Length 22;
81.8%; Pred. No. 1.5e+03;
trive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                         Old L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22 BP; 8 A; 7 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                         Knuth A,
                                                                                                                                               (LUDW-) LUDWIG INST CANCER RES.
(SLOK ) SLOAN KETTERING INST CANCER RES.
(CORR ) CORNELL RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2608 CAAAGCTGAGCCTGCAGGGAAG 2629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 12; Page 16; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAAAGCAGAGCCTCCCGAGAAG 22
                                                                                                                                                                                                                                                                         Scanlan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Snaidr J;
29-NOV-2000; 2000WO-US042334.
                                                         30-NOV-1999; 99US-00451739.
24-OCT-2000; 2000US-00602362.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI68502 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18; Conservative
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                                                                                                                                                                                                                                                                         Stockert B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Walcher M, Wagner M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                      WPI; 2001-441706/47.
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                                                                                                                                               This invention describes a novel method for specifically detecting microorganisms in a sample by Polymerase Chain Reaction (PCR) where in addition to reaction primers specific to the target organism, competition primers specific for non-target organisms are also used. The invention is used to detect microorganisms, particularly to detect infection by Listeria below the species level, especially Listeria monocytogenes. The invention allows detection of different subspecies of Listeria not provided by prior art. This sequence represents a PCR primer used in the amplification of the Listeria monocycogenes as per primer used in the pso protein described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; NOVX; human disease; NOVX-associated disorder; cancer; addiction; Hodgkin disease; Von Hippel-Lindau syndrome; Alzheimer's disease; stroke; tuberous seleroais; hypercalcaemia; Parkinson's disease; depression; tuberous seleroais; parcial palsy; epilepsy; Lesch-Nyhan syndrome; multiple sclerosis; ataxia-telangiectasia; leukodystrophy; anxiety; pain; obesity; Crohn's disease; osteoporosis; inflammatory bowel disease; inferdility; inflammatory bowel disease; scleroderma; haemophilia; diabetes; pancreatitis; autoimmune disease; scleroderma; haemophilia; diabetes; pancreatitis; autoimmune disease; bacterial infection; parasitic infection; graft-versus-host disease; ecell differentiation; cell proliferation; haematopolesis; wound healing; angiogenesis; PCR; primer; ss.
                                        chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                    Specifically detecting microorganisms in a sample, by polymerase chareaction with reaction and competitor primers, useful for detecting subspecies of Listeria, in particular Listeria monocytogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 15.6; DB 1; Length 22; 81.8%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 22 BP; 1 A; 1 C; 7 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2328 TGTGTGCGTGTGTGTGTGTG 2349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 rerefrecterrierarriere 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human NOVX forward primer Ag3477.
                                                                                                                   Claim 10; Page 16; 32pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAR-2001; 2001US-0274849P.

12-MAR-2001; 2001US-0275235P.

13-MAR-2001; 2001US-0275579P.

13-MAR-2001; 2001US-0275601P.

14-MAR-2001; 2001US-0276000P.

20-MAR-2001; 2001US-0277239P.

20-MAR-2001; 2001US-027733P.

21-MAR-2001; 2001US-0277338P.

22-MAR-2001; 2001US-0277331P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABS78766 standard; DNA; 22 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
WPI; 2001-625966/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   #0200272770-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABS78766;
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The present invention relates to a new polypeptide (NOVX). The NOVX polypeptide, nucleic acid and antibody are useful in the manufacture of a medicament for treating a syndrome associated with a human disease, preferably a NOVX associated disorder. The NOVX nucleic acids, polypeptides and antibodies are useful for treating, preventing or diagnosing diseases such as cancers, Hodgkin disease, Von Hippel-Lindau syndrome, Alzheimer's disease, stroke, tuberous sclerosis, crebral palsy, epilepsy, Lesch-Nyhan syndrome, multiple sclerosis, ataxia-clangiectasis, leukodystrophies, addiction, anxiety, depression, pain, obesity, Crohn's disease, osteoporosis, inflammatory bowel disease, infertility, inflammatory bowel disease, atherosclerosis, hypertension, coleroforma, haemophilia, diabetes, pancreatitis, autoimmune disease, athem, arthritis, immunodeficiencies, Huving, viral, bacterial or parasitic infections, or graft-versus-host disease. The nucleic acids and consider that manual as targets for the identification of small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   molecules that modulate or inhibit ergon neurogenesis, cell differentiation, cell proliferation, haematopoiesis, wound healing and anglogenesis, in gene therapy, in generation of antibodies that bind immunospecifically to NOVX subsetances for use in therapeutic or diagnostic methods. The nucleic acids are further used as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine, and pharmacogenomics. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention for amplification of
                                                                                                                                                                                                                                                                                                                                                                                                                   New NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. diabetes, multiple sclerosis, atherosclerosis, cancer, infections, osteoporosis or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                       KA, Vernet CA, Tchernev VT, Malyankar UM, Gerlach VL;
Zerhusen BD, Patturajan M, Gusev VY, Kekuda R, Pena CEA;
J, Gangolli EA, Taupier RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.6; DB 1; Length 22; 11.8%; Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 10 A; 3 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1361 TGAAGATGATCGGGAAACACAA 1382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example C; Page 200; 266pp; English.
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             26-MAR-2001; 2001US-0278894P.
27-MAR-2001; 2001US-0279364P.
30-MAR-2001; 2001US-0279344P.
30-MAR-2001; 2001US-0280802P.
02-MAY-2001; 2001US-0280802P.
31-MAY-2001; 2001US-028114P.
31-OCT-2001; 2001US-0338375P.
04-DEC-2001; 2001US-0338375P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 81.8%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-2002 (first entry)
                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Parkinson's disease.
                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-713508/77.
                                                                                                                                                                                                                                                                                           Spytek KA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABS60863;
                                                                                                                                                                                                                                                                                                                                Zhong M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1156
ABS60863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 田城站城即城田
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08-AUG-2002

polymorphism; haemangioma; tumour; sarcoma; Crohn's disease; trachoma; cardiovascular disease; angina pectoris; hypertennion; heart failure; myocardial infarction; ventricular hypertrophy; vascular disease; aneurysm; embolism; thrombosis; ocronary artery disease; angioedaema; arteriosclerosis; atherosclerosis; hypersensitivity; sepsis; PCR; autoimmune disease; inflammatory arthritis; cancer; wound; genotyping; viral infection; bacterial infection; fungal infection; OPPD; aminopeptidase P; XPNEP2; bradykinin receptor B1; primer; BDKRB1; tachykinin receptor B1; TACR1; C1 esterase inhibitor; CINH; kallikrein 1; KLK1; bradykinin receptor B2; BDKRB2; gene therapy; angiotensin converting enzyme 2; ACB2; protease inhibitor 4; P14; Chronic obstructive pulmonary disease; enterocolitis

Homo sapiens.

WO200261131-A2.

03-DEC-2001; 2001WO-US047235

04-DEC-2000; 2000US-0251015P. 23-JAN-2001; 2001US-0263678P. 02-MAR-2001; 2001US-0273037P.

(BRIM) BRISTOL-MYERS SQUIBB CO. (TSUC/) TSUCHIHASHI Z. (HUIL/) HUI L.

Zerba KE, Ma-Edmonds M, Perrone MH; Tsuchihashi Z, Hui L, Swanson BN, Powell JR;

WPI; 2002-619265/66.

New isolated nucleic acid with at least one polymorphic position, useful for detecting, diagnosing and treating disorders such as angioedema, cancer, viral, bacterial or fungal infection, cardiovascular and autoimmune diseases.

Example 3; Page 891; 977pp; English.

The invention relates to an isolated nucleic acid from a human gene encoding aminopeptidase P (XPNEP2), bradykinin receptor B1 (TAREB1), C1 esterase inhibitor (CLNH), kallikrein taceptor B1 (TAREB1), C1 esterase inhibitor, (CLNH), kallikrein c1 (KLKI), bradykinin receptor B2 (BDKRB2), angiotensin converting enzyme (C 2 (ACE2) or protease inhibitor 4 (P14), comprising at least one prayme coolymorphic position as provided in the detailed summary of single nucleotide polymorphisms comprising additional 5' and 3' flanking genomic sequence; (2) analysing (M1) at least one nucleic acid sample comprising obtaining the sample from one or more polymorphic positions in a gene cocding a protein selected from the group above; (3) constructing (M2) encoding a protein selected from the group above; (3) constructing (M2) haplotypes using the genes comprising grouping at least two nucleic acids (3) identifying (M3) an individual at risk of developing a disorder upon administration of an ACE inhibitor and/or vasopeptidase inhibitor using the polymorphic detai, (5) allibrary of nucleic acids each of which comprises one or more polymorphic positions within a gene encoding a nucleic acid sample, determining the numan protein selected from the group above; and (6) genotyping (M4) an individual comprising obtaining a nucleic acid sample, determining the nucleotide present in at least one polymorphic position, are useful for detecting, diagnosing, treating, there are useful for detecting, diagnosing, treating, involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's involve angiogenesis like haemangiomas, tumours, sarcomas, Crohn's hypertrophy, vascular diseases, and/or atherosclerosis, and/or atherosclerosis, and companing arthritis, cancer, wounds, viral, bacterial or fungal infection, Chronic obstructive pulmonary disease, arteriosclerosis, and/or atherosclerosi; inflammatory disease, contends of the contends of the contends of the contends of the contend

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           polynucleotides are also useful for chromosome identification. Antibodies against the proteins may be utilised for immunophenotyping of cell lines and biological samples. The present sequence is a genotyping PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Composition for treating mammary tumors associated with aberrant tyrosine kinase receptor activity in nonhuman animals, comprises one or more substances that inhibit the aberrant activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumours associated with aberrant tyrosine kinase receptor activity in nonhuman animals. (C) comprises one or more substances that inhibit the aberrant tyrosine kinase receptor activity. (C) has cytostatic activity. (C) can be used as a tyrosine kinase receptor inhibitor, and an epidermal growth factor receptor (BGFR) inhibitor. (C) is especially useful for tracting canine mammary tumours. The present sequence represents a PCR primer for human HBR2 receptor, which is used in an example from the present invention for cloning the canine HBR2 receptor
                                                and biological samples. The present sequence is a genotyping PCR primer for the gene encoding one of the proteins listed above
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a composition (C) for treating mammary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bpidermal growth factor receptor; tyrosine kinase receptor inhibitor;
epidermal growth factor receptor inhibitor; EGFR; mammary tumour;
cytostatic; PCR primer; ss.
                                                                                                                                                                                      Gaps
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diseases and disorders are listed in the specification). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 15.6; DB 1; Length 22;
81.8%; Pred. No. 1.5e+03;
tive 0; Mismatches 4; Indels
                                                                                                                                               Score 15.6; DB 1; Length 22;
Pred. No. 1.5e+03;
                                                                                                                                                                                    4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bette P,
                                                                                                            Seguence 22 BP; 3 A; 1 C; 10 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 22 BP; 4 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human HER2 receptor PCR primer SEQ ID NO:11.
                                                                                                                                                             81.8%; Pred. ...
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Van Meel J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BOEH ) BOEHRINGER INGELHEIM INT GMBH.
                                                                                                                                                                                                                         2326 TGTGTGTGCGTGTGTGTGTG 2347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1748 TGAAGTGGATGGCGCCTGAGGC 1769
                                                                                                                                                                                                                                               1 TGTGTGCATGAGTGTAGGTG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 22; 37pp; English.
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                                                                                                                                                 0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hilberg F, Brandstetter I,
                                                                                                                                                                                                                                                                                                                                                          ACC69082 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                    Matches 18; Conservative
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Best Local Similarity 81.8
Matches 18; Conservative
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                                                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                             ACC69082;
                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                     1157
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Human PDE4A-MTA-derived oligonucleotide SEQ ID 14275.

(first entry)

29-JUL-2004

ABD32064;

BP.

ABD32064 standard; DNA; 22

RESULT 1159 ABD32064/c

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entitlenamatory steroid and ubiquinone. A composition of the invention has antiinflammatory steroid and ubiquinone. A composition of the invention immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing or depleting levels of, or reducing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                   antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Katz E, Pabalan J, Aguilar D;
                                                                                                                                                                                                                                  dysfunction; nasal airway dysfunction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 5 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 14275; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                       lung inflammation; respiratory disease; ds.
                                                                                                                                                                                    Human PDE4A-MTA oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Li Y, Sandrasagra A, Ke
Tang L, Shahabuddin S;
                                              ABZ99033 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                      (first entry)
                                                                                                                                                                                                                                antisense; lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                        17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miller S,
                                                                                       ABZ99033;
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RESULT 1158
                        ABZ99033/c
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comprising oligomucleotides, a novel composition, all state active agent, comprising oligomucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and controlly depletion or hyposecretion, when administered to a mammal. The coligomucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a corresponding a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cult and isease or conditions and/or surfactant hypoproduction are associated with a disease or conditions and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated continue maphysema, chronic obstructive pulmonary disease, pulmonary continues present in the target RNA serves to prevent the breakdown of the inflammation rejection, pulmonary infections, bronchities or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the inflammation and advanced the anti-sense oligos corresponding in the target RNA serves to prevent the breakdown of the particular and an an and an an and an an and an and an an and an and an an an an analysis and an analysis and an a
                                                                                                                                                                                                                             respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory diseases syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the oligonucleotides into products that free adenosine into the system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ç
                                                                                                                                                                                                    antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense obligonuclectide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Katz E, Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 14275; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Li Y, Sandrasagra A, Ke
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miller S,
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Gaps

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0.4%; Score 15.6; DB 1; Length 22; 11.8%; Pred. No. 1.5e+03; ve 0; Mismatches 4; Indels

81.8%;

Best Local Similarity

Query Match

18;

Matches

Conservative

1879 GAGCTCTTCAAGCTGCTGAAGG 1900

GTGGGCTTCAAGCTGCTGCAGG 1

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ADG82621 standard; DNA; 22

RESULT 1161

ADG82621

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The present invention describes a method (M1) for estimating therapeutic efficacy of tumour necrosis factor (TNF). M1 involves evaluating the expression level of a TNF-related gene in a cancer cell. Also described is a kit for estimating the therapeutic efficacy of TNF, which is used in the treatment of cancers. The kit comprises a thermostable DNA polymerase and an oligonucleotide primer comprising a DNA sequence encoding a gene chosen from a protein kinase B (Akt-1) gene, death receptor (DR3) gene, multidrug resistance-associated protein (MRP5) gene, and multidrug resistance-associated protein (MRP5) gene, meather sequence represents a PCR primer which is used in an example from the present
e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour necrosis factor; TNF; tumour necrosis factor alpha; TNF-alpha; TNF-related gene; TNF-alpha-related gene; cancer; human; FCR primer; ss.
                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Estimating therapeutic efficacy of tumor necrosis factor involves evaluating expression level of tumor necrosis factor-related gene in
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                                                                                         / Match 0.4%; Score 15.6; DB 1; Length 22; Local Similarity 81.8%; Pred. No. 1.5e+03; nes 18; Conservative 0; Mismatches 4; Indels
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81.8%; Pred. No. 1.5e+03;
ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           TNF-alpha-related gene ERK1 PCR primer SEQ ID NO:55.
                                                     Sequence 22 BP; 5 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 22 BP; 6 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yanai Y, Yamamoto S, Yamamoto K, Ikegami H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; SEQ ID NO 55; 56pp; English.
                   prevent any unwanted effects due to it
                                                                                                                                                                    1879 GAGCTCTTCAAGCTGCTGAAGG 1900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HAYB ) HAYASHIBARA SEIBUTSU KAGAKU
                                                                                                                                                                                                        GTGGGCTTCAAGCTGCTGCAGG 1
                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-2002; 2002JP-00107126.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-APR-2003; 2003EP-00252225.
                                                                                                                                                                                                                                                                                              ADG09487 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer cell.
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                                                                                                                                                                                                                                                                                                                                    ADG09487;
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                                                                                           Query Match
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                                                                                                                               Matches
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The present invention describes a method for promoting (MI) liver growth or promoting (M2) hepatocyte proliferation in the liver of a subject, cor promoting (M3) a pathological liver condition in a subject, or protecting (M4) liver from damage in the subject due to exposure to a hepatotoxic agent, which involves administering to the subject a vascular endothelial crowd in the factor receptor (VBGFR) modulating agent (I). Also described, (I) an article of manufacture comprising a container, composition comprisies of the composition for promoting liver growth, where the composition comprises of the composition comprisies a liver growth, where the composition comprises a vEGFR modulating agent in the amount effective to promote liver growth, a second container on the first container, and a composition comprises a VEGFR modulating agent in the amount cffective to promote liver growth, a second container comprising a buffer of the composition comprises a VEGFR modulating agent in the amount of the composition for using the kit for promoting liver growth, a second container and an instruction for using the kit for promoting liver growth. (I) has hepatotropic and antiliflammatory activities, and can be used for promoting liver growth promoter. (I) can be used for promoting crowth or hepatocyte proliferation in a subject such as liver creating a pathological liver form damage in a subject such as liver from damage in a subject come or hepatotoxic agent. The VEGFR modulator create a local cascade of signaling events originating in sinusoidal endothelial cells following to vEGFR receptor activation, which is much more potent and beneficial in promoting hepatocyte proliferation and liver growth than systemic containers sequence is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Promoting liver growth or promoting hepatocyte proliferation in liver of subject, treating pathological liver condition e.g. cirrhosis in subject, by administering vascular endothelial growth factor receptor modulator.
                                                                                                                                                    liver growth, hepatocyte proliferation, pathological liver condition, liver damage; vascular endothelial growth factor receptor modulator; VEGFR modulator; hepatitispion antiinflammatory; liver growth promoter; liver failure; hepatitis; liver cirrhosis; toxic liver damage; medicamentary liver damage; hepatic encephalopathy; hepatic coma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 6 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Le Couter J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 4; Page 43; 64pp; English.
                                                                                                                                                                                                                                                                            hepatic necrosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUN-2003; 2003WO-US017591.
                                                                                                            bFGF gene forward PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUN-2002; 2002US-0386637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hillan KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-071254/07.
                                                                                                                                                                                                                                                                                                                                                                   WO2003103581-A2.
                                                                11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2003,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ferrara N,
                   ADG82621;
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Score 15.6; DB 1; Length 22; Pred. No. 1.5e+03;

0.4%; 81.8%;

Query Match Best Local Similarity

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Gaps

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Local Similarity 81.8%; les 18; Conservative

Best Loca Matches

(first entry)

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interleukin, IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                            Oligonucleotide associated to PDE4C #351.
                                                                                                                                                                                                                                                                                                                                                                                                    Nyce JW, Tang L, Sandrasagra A,
Shahabuddin S, Lu H, Cong H;
                            ADJ61285 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                             29-JUL-2002; 2002US-0399076P.
                                                                                                                                                                                                                                                                                                                25-JUL-2003; 2003WO-US023509.
                                                                                                                                                                                                                                                                                                                                                                        (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-203534/19.
                                                                                                                                                                                                                                                         WO2004011613-A2
                                                                                                                                                                                                                              Homo sapiens.
                                                                                 06-MAY-2004
                                                                                                                                                                                                                                                                                   05-FEB-2004
                                                      ADJ61285;
RESULT 1163
ADJ61285/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method for estimating the therapeutic effect of interferon in the treatment of cancer or viral disease. The method gene involves determining the amount of expression of an interferon-associated gene in cancer cells or virus-infected cells. The invention also relates to drug compositions for the treatment of cancer and viral diseases containing interferon-alpha together with a cycloxygenase-2 (Cox.2) inhibitor such as indomethacin which potentiates the apoptosis induction are useful in the treatment and prevention of cancers (e.g., cancer of the colon, lung, pancreas, breast, stomach, liver, kidney, nerve cell, skin, muscle, uterus and throat) and viral infections (e.g., hepatitis B and C). The present sequence represents a PCR primer used in real-time PCR to determine the amount of expression of an interferon-associated
                                                                                                                                                                                                                                        Interferon therapy, cancer, viral disease, viral infection, interferon-alpha; IFN-alpha; cyclooxygenase-2 inhibitor; Cox-2 inhibitor; apoptosis induction; colon cancer; lung cancer; pancreas cancer; presast cancer; promach cancer; liver cancer; kidney cancer; nerve cell cancer; skin cancer; muscle cancer; uterus cancer; throat cancer; hepatitis B; hepatitis C; cytostatic; virucide; cancer cell; interferon-associated gene; ERK1; real-time PCR; primer; ss.
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Measurement of Cox-2 gene expression in cancer or virus-infected cell for estimating the therapeutic effect of an interferon in cancer and viral disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and C). The present sequence any procession of an interior and PCR to determine the amount of expression of an interferon-alpha.
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4; Indels
                                                                                                                                                                                                              IFN-associated gene ERK1 PCR primer, SEQ ID NO:55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 6 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yamauchi H;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 55; 90pp; Japanese.
                            1979 CCTCCCAGAGGCCCACCTTCAA 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HAYB ) HAYASHIBARA SEIBUTSU KAGAKU.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yamamoto K,
                                           1 CCTCTCAGAGACCTACGTTCAA 22
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                                                                                                                            BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-JUL-2002; 2002JP-00195147
                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JUN-2003; 2003WO-JP008296
                                                                                                                            ADH75266 standard; DNA; 22
                                                                                                                                                                                   22-APR-2004 (first entry)
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yamamoto S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-108824/11.
                                                                                                                                                                                                                                                                                                                                                                                       WO2004005549-A1
                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                    15-JAN-2004.
 18;
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                                                                                                                                                        ADH75266;
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                                                                                              Matches
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Aguilar D, Miller S;

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                                                   e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              יאמטאי, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant gener CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 15.6; DB 1; Length 22; 81.8%; Pred. No. 1.5e+03; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 5 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                   Claim 2; SEQ ID NO 2141; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1879 GAGCTCTTCAAGCTGCTGAAGG 1900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gradactricaactractacass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ60916 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ60916;
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ADJ60916/c
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855 GGAGGAGCTGGTGGAGGCTGAC 876

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18; Conservative

Local Similarity

Best Loca Matches

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The present invention relates to an oligonuclectide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL) 4 receptor, IL-5 receptor or salts of the oligonuclectide and optionally surfactant operatively linked to the oligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease. CC chosen from airway inflammation, allergy(iss), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CRDD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway contaction. The present sequence represents an oligonucleotide of the
                                                                                                                     airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        initiation codons and introns of respiratory disease-relevant gener
CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel single or multiple target oligonucleotide anti-sense to
                                                                                                 interleukin; IL-4 receptor; IL-5 receptor; lung disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nyce JW, Tang L, Sandrasagra A, Aguilar D, Miller S; Shahabuddin S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 5 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                            Oligonucleotide associated to PDE4A #199.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; SEQ ID NO 1772; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                       29-JUL-2002; 2002US-0399076P.
                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                               25-JUL-2003; 2003WO-US023509
                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                           WO2004011613-A2
                                                                                                                                                                                                                                  Homo sapiens
                    06-MAY-2004
                                                                                                                                                                                                                                                                                                                     05-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention.
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Gaps ö Score 15.6; DB 1; Length 22; Pred. No. 1.5e+03; 4; Indels 0; Mismatches ch 1 Similarity 81.8%; 18; Conservative Query Match Best Local Similarity Matches 18; Conserv

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1879 GAGCTCTTCAAGCTGCTGAAGG 1900 22 grasscricaascriscrass

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Human oligonucleotide #1771. (first entry) 15-JUL-2004 ADO46405; AD046405/ ID AD04 XX AD04 XX AD04 XX DT 15-J XX XX

ADO46405 standard; DNA;

RESULT 1165

lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory diseases; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction. IL-4; interleukin-5 receptor; IL-5; Human, 85; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;

Homo sapiens.

US2004049022-A1

11-MAR-2004

25-JUL-2003; 2003US-00627930.

23-APR-2002; 2002WO-US013135. 23-APR-2002; 2002WO-US013143.

NYCE/)

NYCE J W. SANDRASAGRA A. SAND/)

TANG L. TANG/)

AGUILAR D. MILLER S. SHAHABUDDIN S. AGUI/)

(MILL/) (SHAH/) (LUHH/)

e.g. genes e.g.,

Aguilar D, Miller S; Sandrasagra A, Tang L, n S, Lu H, Cong H; CONG H. Š CONG/) Nyce

Shahabuddin S, Lu H, WPI; 2004-293804/27. Novel single or multiple target oligonuclectide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.

Claim 2; SEQ ID NO 1772; 174pp; English.

The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Edtakin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Edtakin PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(S) or expressed product(S), for the prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a conjunct ociding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Edtakin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for reating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A inflammation or an inflammation, allergies associated with inflammation or an inflammation, allergy, asthma, impeded respiration, cystific fibrosis (CP), chronic disease. The respiratory or lung disease is associated with impeded respiration, cystific fibrosis (CP), chronic obstructive pulmonary disease. allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the invention

Sequence 22 BP; 5 A; 9 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö Score 15.6; DB 1; Length 22; Pred. No. 1.5e+03; 0; Mismatches 4; Indels 0.4%; ilarity 81.8%; Conservative Query Match Best Local Similarity Matches 18; Conserv

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receptor(s), and/or asthma and/or lung allergies associated with inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary disease (COPD), allergic fibrosis (CP), chronic obstructive pulmonary disease (COPD), hypertension, bung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the

Sequence 22 BP; 5 A; 9 C; 5 G; 3 T; 0 U; 0 Other;

invention.

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Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                  CCRI; CCR3; ECEATION.

CCRI; CCR3; ECEATION.

CCR1; CCR3; ECEATION.

CCR1; CCR3; ECEATION.

CCR1; CCR3; ECEATION.

Lung disease; hyper-responsivenes; adenosine, adenosine, acceptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; inflammatory disease; airway inflammation; dispraction; allergy; is a sease; copp. allergy; allergy; inspedd respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CCP9; allergic fibritis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction,
                                                                                                                                                                         interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Miller S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aguilar D,
1879 GAGCTCTTCAAGCTGCTGAAGG 1900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tang L,
ng H;
                       22 GTGGGCTTCAAGCTGCTGCAGG 1
                                                                               ADO46675 Btandard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                         23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                  25-JUL-2003; 2003US-00627930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sandrasagra A, Ta
                                                                                                                                                   Human oligonucleotide #2041,
                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                           NYCE J W.
SANDRASAGRA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MILLER S.
SHAHABUDDIN S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TANG L.
AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-293804/27
                                                                                                                                                                                                                                                                                                                     JS2004049022-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shahabuddin S,
                                                                                                                             15-JUL-2004
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B
                                                                                                      ADO46675;
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(TANG/)
(AGUI/)
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                                                         RESULT 1166
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The present invention relates to a method of providing neuroprotection in a subject, which comprises administering to the subject a ciliary curvothic factor (CMTP) peptide or its biologically-active fragment or variant and a peptide that acts as a receptor for active fragment or variant and a peptide that acts as a receptor for CMTP. The method, composition and compound are useful in providing neuroprotection in a subject. The peptide having the activity of a CMTP peptide and the peptide that acts as a receptor for CMTP are useful in providing neuroprotection. For modulating cell degeneration or cell death, for treating or preventing a medicament for neuroprotection, for modulating cell compound. The method, composition and compound are useful in treating or preventing a subject with substance abuse, trauma, stroke, ischaemia, Huntington's disease, Alzheimer's disease, Parkinson's disease, prion disease, variant Creutzfeld-dakeb disease, amyotrophic lateral scleresis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ALS), olivopontocerebellar atrophy, epilepsy, seizures or hypoglycaemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Providing neuroprotection in a subject comprises administering to the subject a ciliary neurotrophic factor (CNTF) peptide or a peptide that acts as a receptor for CNTF peptide.
                                          Gaps
                                                                                                                                                                                                                                                                                                                                                         cerebroprotective; vasotropic; anticonvulsant; nootropic;
neuroprotective; antiparkinsonian; antiapoptotic; antiaddictive;
vulnerary; tranquilizer; CNTF; ciliary neurotrophic factor;
                                        ö
Score 15.6; DB 1; Length 22;
Pred. No. 1.5e+03;
0; Mismatches 4; Indels
                                                                           1879 GAGCTCTTCAAGCTGCTGAAGG 1900
                                                                                                                22 Gregerrchagerecrecaes 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example A; Page 23; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                       neuroprotection, PCR; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYBR-) UNIV BRITISH COLUMBIA.
                                                                                                                                                                                                             ВР.
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   0.4%;
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                                                                                                                                                                                                             ADO05994 standard; DNA; 22
                                                                                                                                                                                                                                                                                   29-JUL-2004 (first entry)
Query Match
Best Local Similarity 81.8
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bechberger J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-365450/34.
                                                                                                                                                                                                                                                                                                                          Cx43 PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2004037281-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAY-2004.
                                                                                                                                                                                                                                                  AD005994;
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                                                                                                                                                                        RESULT 1167
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 mucleotides of the 5'-end or 3'-end of a mucleic acid target consen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The coligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are treptratory or lung disease. The respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A

Claim 2; SEQ ID NO 2141; 174pp; English.

asthma.

HIV; thiolation; reverse transcriptase; primer; inhibition; homooligomer;

Anti-sense oligonucleotide specific for tRNA_Lys anti-codon region

(first entry)

13-JUL-1992

AAQ25023

30 ATATATATAAAAAAAAAAAAAAAAA 1

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AAQ25023 standard; DNA; 40

RESULT 1169

4AQ25023

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encelopathy. The present sequence is a PCR primer used to isolate a Cx43 coding sequence in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Probe; detection; genotyping; cell status; chromosomal resistance gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting specific interactions between molecular targets and probes, useful for determining genotypic and physiological status of cells, by attaching an markable adapter to the target.
                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel method for detecting the specific
                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 15.6; DB 1; Length 30; 70.0%; Pred. No. 2e+03; tive 0; Mismatches 9; Indels
                                                                                                                                                                                                                                                 4; Indels
                                                                                                                                                                         Score 15.6; DB 1; Length Pred. No. 1.5e+03; 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wagenhaus A, Ermantraut E;
                                                                                                     Sequence 22 BP; 3 A; 12 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 30 BP; 5 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CLON-) CLONDIAG CHIP TECHNOLOGIES GMBH
                                                                                                                                                                                                                                                                                                               1119 CCCCACGCTGGCCAATGTCTCC 1140
                                                                                                                                                                                                                                                                                                                                                                                cccacrcrcaccrarererec 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 8; 25pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                         Query Match
Best Local Similarity 81.8%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-APR-2001; 2001DE-01017857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-APR-2001; 2001DE-01017857,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX89953 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             described in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 70.0°
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ellinger T, Ehricht R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-068801/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PolyA adapter DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DE10117857-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABX89953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   array; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABX89953/c
ABX89953/c
ABX899953/X
AC ABX899955
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Compsn. contg. 5-thiolated (oligo-poly-)-nucleotide(s) - for treating HIV infection, AIDS and for preventing HIV-1 infection.

Disclosure; Page 12; 42pp; English

Schinazi RF;

Bardos TJ, Ho YK, Aradi J,

WPI; 1992-096567/12.

(UYNY-) RES FOUND UNIV NEW

91WO-US005919. 90US-00568131,

15-AUG-1991; 16-AUG-1990;

WO9203127-A 05-MAR-1992

Synthetic.

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                                                                                                                                                                                                                                                                                                           The oligomer comprises an 18-mer regional oligonucleotide corrresponding to the anti-codon region of the reverse transcriptase (RT) primer tRNA_Lys, with a S' terminal thiolated 22-mer homooligomer. The homooligomer may be absent and the regional oligo may itself be thiolated (e.g. 5-Mercapto-), as in Claim 19. The oligos are used to inhibit reverse transcription in the treatment of HIV. See also AAQ22624-27 and
                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                    ô
                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 20.0%; Pred. No. 2.4e+03; 6; Conservative 15; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40 BP; 7 A; 2 C; 3 G; 6 T; 22 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         3264 TTTTATTTGCTTTGTCCTTTTTCAGGAGAA 3293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               202P5A5; human; cancer; tumour; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human 202P5A5 cDNA synthesis primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADN89686 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004016736-A2.
                                                                                                                                                                                                                                                                                                                                                                        AAQ25017-Q25032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADN89686;
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2834 ATATATATATAACATATATGGAAGGGAAA 2863

26-FEB-2004.

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Gaps

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                                    This sequence represents a primer used in the isolation of cDNA fragments of the PTAM (testis specific protein expressed in prostate cancer) gene. PTAM is expressed in 1 isoforms PTAM-1, 2, and 3. The PTAM gene is located on chromosome 1922. PTAM is overexpressed in prostate cancer, and has a testis specific expression pattern in adult tissues. PTAM shows no homology to any known gene. PTAM can be used in methods for the diagnosis of cancer, especially prostate or breast cancer, where the normal tissue samples are prostate tissue, or breast tissue, bone tissue, lymphatic tissue, a vaccine composition targeting PTAM, PTAM, ribozymes specific for PTAM and antisense sequences, can be used to treat cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention comprises a composition which contains a substance that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3
                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122P5C8; cancer-associated gene; cancer-associated protein; cancer; cDNA synthesis; primer; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New composition comprising a substance that modulates the status of 125P5C8 gene or a molecule that is modulated by 125P5C8, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g
                                                                                                                                                                                                  Cancer development can be
                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Raitano AB,
                                                                                                                                                                                                                                                                          Score 15.6; DB 1; Length 42; Pred. No. 2.4e+03;
                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Afar DEH,
A,
                                                                                                                                                                                                                                               Sequence 42 BP; 3 A; 2 C; 2 G; 35 T; 0 U; 0 Other;
                                                                                                                                                                                                    especially breast and prostate cancers. Cancer devinhibited by a vaccine composition targeting PTAN
                                                                                                                                                                                                                                                                                                         6
                                                                                                                                                                                                                                                                                                                                    3256 TGAAGATATTTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              125P5C8 gene-related cDNA synthesis primer.
                                                                                                                                                                                                                                                                                                                                                                 6 rcaagcririririririririririririri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 714; 183pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hubert RS,
Jakobovits
             Example 1; Page 31; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosing or treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                         ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-MAR-2002; 2002US-00099460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAR-2001; 2001US-00809638
                                                                                                                                                                                                                                                                            0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KJM,
                                                                                                                                                                                                                                                                                                                                                                                                                                         ADL19444 standard; DNA; 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 70.0°
Matches 21, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MORRISON R K.
MORRISON K J M.
JAKOBOVITS A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FARIS M.
CHALLITA-EID P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Challita-Eid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Morrison RK, Morrison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUBERT R S.
AFAR D E H.
RAITANO A B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-021932/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003219444-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADL19444;
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(MORR/) I
(JAKO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Faris M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FARI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHAL/)
(HUBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RAIT/)
(GEWW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AFAR/)
                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1172
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                                                                                                                                                                       New compositions having the 202P5A5 gene and encoded protein, useful for diagnosing, preventing, prognosticating or treating cancer of the prostate, bladder, colon, lung, ovary, breast, stomach, cervix, lymphoma,
                                                                                                                                                                                                                                                                          The invention relates to a composition comprising 202P5A5 proteins. The composition and proteins are useful for detecting and treating cancer by inhibiting the growth or viability of cancer cells. The present sequence represents the human 202P5A5 cDNA synthesis primer.
                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer; overexpress; chromosome 1q22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PTAN proteins, and sequences encoding them, used for diagnosing and treating cancers, especially breast and prostate cancers.
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                                                                                                                                                                                                                                                                                                                                                                               Score 15.6; DB 1; Length 41;
Pred. No. 2.4e+03;
0; Mismatches 9; Indels
                                                                                                                  e
                                                                                                                Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                  Sequence 41 BP; 3 A; 2 C; 2 G; 34 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA synthesis primer used in PTAN gene isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PTAN; testis specific; prostate cancer; overe: diagnose; cancer; breast; vaccine; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                         3256 TGAAGATATTTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mitchell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rcaagcirririrrirrirrirrirrirririr 36
                                                                                                               Challita-Eid PM,
                                                                                                                                                                                                                                                Example 1; SEQ ID NO 28; 266pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Raitano AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA37946 standard; DNA; 42 BP.
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98US-0113229P.
99US-0129518P.
                                                                                                                                                                                                                                                                                                                                                                             0.4%;
             16-JUN-2003; 2003WO-US018906.
                                       16-AUG-2002; 2002US-0404306P
01-NOV-2002; 2002US-0423290P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 70.0°
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UROGENESYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AFAR D E.
HUBERT R S.
RAITANO A B.
MITCHELL S C.
                                                                                                               Raitano AB, Faris M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hubert RS,
                                                                                                                                            WPI; 2004-203774/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-317715/27
                                                                                   (AGEN-) AGENSYS INC
                                                                                                                                                                                                                   bone and/or skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200020589-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-OCT-1998;
21-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UROG-)
(AFAR/)
(HUBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Afar DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RAIT/)
(MITC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1171
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ઠે 셤 70.0%;

Best Local Similarity

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The invention relates to a method of determining the presence or absence of a CYP2D6 target sequence in a DNA sample containing CYP2D6 nucleic acid. Determining the presence or absence of a CYP2D6 target sequence in a sample of DNA containing CYP2D6 nucleic acid comprises contacting the nucleic acid with a probe under stringent binding conditions, and detecting the presence or absence of the target sequence bound with the probe with a scattered light detectable particle, by observing light scattered from the particle which indicates the presence of the target sequence. The method is useful for determining the presence of absence of particular single nucleotide polymorphisms or alleles in genomic nucleic acid, especially in a pharmacogenetically relevant gene or genes in a DNA sample, and to detect and measure one or more target sequences in a sample, the method may also be used to detect specific mutations to identify the phenotypic classification of an individual. ABX30162-
modulate the status of 125P5C8 (125P5C8 is a novel cancer-associated gene/protein), or a molecule that is modulated by 125P5C8. The composition of the invention is useful for diagnosing or treating cancer. The present DNA sequence represents a cDNA synthesis primer that was used in an example of the invention to isolate a fragment of the 125P5C8 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Determining the presence of a CYP2D6 target sequence in a DNA sample containing CYP2D6 nucleic acid, for detecting mutations or polymorphisms, comprises detecting the scattered light from a particle bound to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; CYP2D6; primer; single nucleotide polymorphism detection; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                            0.4%; Score 15.6; DB 1; Length 42; 70.0%; Pred. No. 2.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yguerabide J;
                                                                                                                                                                                                                        Sequence 42 BP; 3 A; 2 C; 2 G; 35 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CYP2D6 gene polymorphism detection primer #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                      3256 TGAAGATATTTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rcaascriririririririririririririririr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peterson T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Fig 6; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK30223 standard; DNA; 48 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUN-2000; 2000US-0210988P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Korb L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                             Local Similarity 70.0
Hes 21; Conservative
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                                                                                                                                                                                                                                                                                            Query Match
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DB 1; Length 48;

Score 15.6;

0.4%;

Query Match

Sequence 48 BP; 33 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

invention

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a segment of the hepatoma AS-30D Type II blackhinase promoter region. Response elements (transcription factor binding site) in this fragment may consist of all or part of the present sequence. AS-30D is a new isolated hexokinase II. The present The present reading frame and contains at least one response element. The present DNA fragment may be coupled to a reporter gene and used to screen for potential drugs that affect regulated transcription of tumour hexokinase II. Alternatively it may be coupled to a toxic gene and used to treat cells that over-express hexokinase II, such as those present in patients with cancer. It may also be used in gene therapy to treat diabetes. The heterologous protein. Probes of the DNA fragment are used in the method for diagnosing a neoplasia that over-expresses hexokinase. The new for diagnosing a neoplasia that over-expresses hexokinase. The new
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New transcription regulating fragments of hexokinase II DNA contg. response element - and methods for diagnosis or treatment of neoplasias that over-express hexokinase II and for regulating glycolysis.
                                                                                                                                                                                                                                                                                                                      Response element; Z-DNA; neoplasia; hexokinase II; glycolysis; cancer;
                                   Gaps
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                                                                                                                                                                                                                                                                                     Hepatoma AS-30D Type II hexokinase promoter fragment from -3843.
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Pred. No. 2.5e+03;
0; Mismatches 14; Indels
                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 49 BP; 3 A; 5 C; 20 G; 21 T; 0 U; 0 Other;
Pred. No. 2.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CACACAGACGTGTACACACACACACACACACACACA
                               0; Mismatches
                                                                                                   3262 TATTTTATTTGCTTTGTCCTTTTTCAGGAG
                                                                                                                                                                                                                                                                                                                                      gene therapy; diabetes; tumour; rat; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Fig 11; 104pp; English
                                                                                                                                                                                    ВР
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                                                                                                                                                                                    AAT80427 standard; DNA; 49
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                                 21; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                        WO9704104-A2
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AAX73034
ID AAX73034
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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vaccular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX7575 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                           stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.4; DB 1; Length 17;
18.8%; Pred. No. 1.2e+03;
ve 6; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human KDR VEGF receptor hammerhead ribozyme substrate #472.
                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecule modulating VEGF receptor(s) gene
                                                                                                                                                                                                                                                                                                                          Stinchcomb D, Escobedo J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; 3 C; 5 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 112; 218pp; English.
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Best Local Similarity 58.8%;
Matches 10; Conservative
                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
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foetal liver kinase 1; ss.
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                                                                                                                                                                                                                                                                                                                          Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                  (CHIR ) CHIRON CORP.
                                          Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyrosine kinase 1 (fll-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75722 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                       Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                    tyrosine kinase 1; kinase insert domain containing receptor;
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                                                                                                 Mouse flk-1 VEGF receptor hammerhead ribozyme substrate #467.
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                                                                                                                                                                                                                         Foetal liver kinase 1; ss
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les 14; Conservative
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                     AAX73034;
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Gaps

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expression or mRNA

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Synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient prefetors of vascular endothelial growth factor (VEGF). A patient receptors of vascular having a condition associated with the level of the fims-like tyrosine kinase 1 (fll-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour anglogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                      present invention describes nucleic acid molecules which modulate the
                                                                                                                                                                                        Nucleic acid molecule modulating VEGF receptor(s) gene expression or stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 6 A; 3 C; 5 G; 0 T; 3 U; 0 Other;
                                                                                                        Stinchcomb D,
                                                                                                                                                                                                                                                                            Claim 4; Page 111; 218pp; English
96US-00584040
                                     RIBOZYME PHARM INC.
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                                                                                                        Pavco P, Mcswiggen J,
                                                           CHIRON CORP
                                                                                                                                                WPI; 1997-259017/23
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11-JAN-1996;
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expression or mRNA

Escopedo J;

0.4%; Score 15.4; DB 1; Length 17; 82.4%; Pred. No. 1.2e+03; tive 2; Mismatches 1; Indel8 1609 AAGTGCATCCACAGGGA 1625 AAGUGUAUCCACAGGGA 17 ઠ 셤

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Gaps

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Nucleotide sequence of the PCR primer 15. AAV36994 standard; DNA; 17 BP (first entry) 24-SEP-1998 AAV36994; 1178

PCR; primer; amplification; type II diabetes; agonist; antagonist; prepatic nuclear factor 4; HNF-4; HNF gene; anti-HNF antibody; insulin; diabetes mellitus; mature onset diabetes of the young; MODY; MODYI gene; MODY3 gene; RS.

Synthetic.

WO9821363-A1

22-MAY-1998

14-NOV-1997;

96US-00749430 15-NOV-1996;

97WO-US020759

(MILL-) MILLENNIUM PHARM INC

Glucksman AM, Thomas J;

WPI; 1998-297964/26.

Treating type II diabetes involving hepatic nuclear factor 4 - useful, e.g. to treat insufficient HNF expression or bioactivity, overexpression of HNF or expression of mutant HNF gene in diabetic patients.

Disclosure, Page 86, 116pp, English

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This is the nucleotide sequence of the PCR primer used for amplification in the method of invention, involving the treatment of type II diabetes with hepatic nuclear factor 4 (HNF-4). The agonists of normal HNF bioactivity can be used to treat diabetes, e.g. to ameliorate disease symptoms involving insufficient expression of an HNF gene and/or inadequate functional HNF bioactivity can be used to treat diabetes, e.g. to ameliorate disease symptoms involving expression of a mutant HNF gene or overexpression of a normal HNF gene. It is also useful to ameliorate disease symptoms involving expression of a mutant HNF gene or overexpression of a normal HNF gene. It is also useful to ameliorate disease symptoms involving a mutant (non-functional) HNF protein e.g. by administering a therapeutically effective amount of an anti-HNF antibody. Protein that binds to the mutant HNF-4 protein is useful in screening assays, e.g. to identify antagonists/agonists of an interaction between a CNF protein and a binding protein to develop drugs for disease treatment. Diabetes mellitus is a common metabolic disorder, and most cases are type of inconinguin dependent diabetes mellitus (NIDDM). A major genetic component is implicated, but few genes have been identified. Mature onset diabetes of the young (MODY) loci have been identified. HNF-4 is encoded by a gene mapping within the MODY1 locus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This is the nucleotide sequence of the PCR primer used for amplification in the method of the invention, which involves modulating the expression of hepatic nuclear factor or other diabetes related gene. The method is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating type II diabetes with agent - useful for, e.g. modulating expression of hepatic nuclear factor or other diabetes-related gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR; primer; amplification; hepatic nuclear factor; HNF; diabetes; type II diabetes; HNF1 gene; transcription factor; insulin; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 2 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.4; DB 1;
Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleotide sequence of 3' PCR primer 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         54 GCTGCAGGTGCTGAATG
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 94.1;
Matches 16; Conservative
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10-JAN-1997;
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The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARWT) gene, an integrin subunit beta 3 gene, an integrin subunit beta 3 gene, an integrin subunit beta 3 corresponding target esquences; and AAA1765 to AAA1767 and AAA17650 and AAA1762 represent ribozyme sequences for ARNT, and AAA1168 to AAA17660 and AAA17685 to AAA18386 to AAA19086 corresponding target sequences; AAA17685 to AAA18386 to AAA19086 and AAA19155 to AAA21922 represent their corresponding target sequences; AAA19155 to AAA21801 and AAA21801 to AAA21805 represent ribozyme sequences; AAA1895 to AAA2160 and AAA21696 to AAA21688 represent their corresponding target sequences; AAA21696 to AAA2168 represent their corresponding target sequences; AAA21699 to AAA22475 and AAA23283 to AAA23362 to AAA21500 and AAA23489 to AAA22435 and AAA23283 to AAA23362, AAA23302, AAA2343 to integrin subunit beta 3, and AAA2346 to AAA2343 the invention are used for modulating the synthesis, expression and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antiniflammatory; antiarthritic; antipsoriatic; ARMD; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma; myopic degeneration; psoriasis; veruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; ss. Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss.
used to treat early onset type II diabetes and defects in insulin secretion. It is based on the discovery that certain mutations in the HNFI gene, encoding a transcription factor, are involved in these conditions
                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                          Match 0.4%; Score 15.4; DB 1; Length 17; Local Similarity 94.1%; Pred. No. 1.2e+03; es 16; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                         Sequence 17 BP; 2 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human TIE-2 substrate sequence SEQ ID NO:2093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 56; Page 122; 305pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                 54 GCTGCAGGTGCTGAATG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GCTGCAGGTGCTGGATG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-591315/50.
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                                                                                                                                                                                                                                                              Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence is a PCR primer used in the cloning of the T66Bk gene. The invention related to a nucleic acid molecule (I) comprising a transcription unit encoding in its 5' portion a kinase having a homology to MARK2 kinase and the 3' portion of the nucleotide sequence has a high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transcription unit; MARK2 kinase; rsk3 kinase; regulatory region; T66Bk;
contraceptive; Responder/Distorter signalling cascade; t-Responder;
stability of an mRNA encoding angiogenic factor, especially ARNT, integrin subunit beta-3, integrin subunit alpha-6, or Tie-2. They are especially used to treat cancer, diabetic retinopathy, age related macular degeneration (ARND), inflammation, and arthritis, as well as neovascular glaucoma, myopic degeneration, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, por wine stains, Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-Rendu syndrome, and other syndromes and diseases related to the levels of ARNT, Tie-2,
                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                          0.4%; Score 15.4; DB 1; Length 17; 12.4%; Pred. No. 1.2e+03; ve 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids involved in the Responder phenotype in mice.
                                                                                                                                                                 integrin subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                     Sequence 17 BP; 3 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer for cloning of T66Bk gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 59; 117pp; English.
                                                                                                                                                                                                                                                                                                                           1619 ACAGGGACCTGGCTGCC 1635
                                                                                                                                                                                                                                                                                                                                             1 ACAGGGAUCUGGCUGCC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
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Best Local Similarity 82.4%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-NOV-1998;
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           also allow distortion, to a non-Mendelian ratio, of the transmission of genetic trait, i.e. determination of sex, from male mammals to their offspring by expressing during spermatogenesis/spermiogenesis a gene involved in sperm motility and/or fertilisation. The genes and proteins involved in the responder phenotype and Responder/Distorter signalling cascade, as well as the inventive methods are advantageous in breeding strategies by allowing for specific selection of genetic traits and in particular, of sex
Responder/Distorter signal cascade other than the t-Responder. They
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detection of chromosome aberrations, used for detecting diseases and disorders, infections, and plant alterations related to e.g. herbicide
                                                                                                                                                                                         Gaps
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                                                                                                                                                           0.4%; Score 15.4; DB 1; Length 17;
94.1%; Pred. No. 1.2e+03;
Ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                            Probe hybridising to nucleotides of human c-erb-B-2 (HER-2)
                                                                                                                                                                                                                                                                                                                                                                                                                     Human, c-erb-B-2; HER-2; chromosome aberration; probe; peptide nucleic acid; haemapoietic malignancy; cancer; inborn constituel disease; herbicide resistance gene; ss.
                                                                                                                                  Sequence 17 BP; 3 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Adelhorst K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 44; 63pp; English.
                                                                                                                                                                                                                  1747 GTGAAGTGGATGGCGCC 1763
                                                                                                                                                                                                                                                                                                               AAZ36590 standard; DNA; 17 BP
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                                                                                                                                                                         94.18;
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                                                                                                                                                                           Local Similarity 94.1
nes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; 88.
                                                                                                                     Gaps
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                                                                               Length 17;
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                                       Sequence 17 BP; 4 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                             Score 15.4; DB 1;
Pred. No. 1.2e+03;
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2001WO-US000662.
2001WO-US000663.
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                                                                             0.4%;
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2000GB-00024263
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                                                                                                                                                                                                     17 AGACTICGGGCTGGCTC
                                                                                                                                                                                                                                                                                                      ABN08004 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                     Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-179446/23.
                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200192524-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                              ABN08004;
                                                                               Query Match
                                                                                                                                                                                                                                                          RESULT 1183
ABN08004
SXS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human c-erb-B-2 (HER-2) gene. The probes are used to demonstrate the method of the invention. The specification describes a method for the method of the invention. The specification describes a method for the detection of chromosome aberrations in eukaryotic samples uses sets of peptide nucleic acid (PNA) probes in hybridisation reactions. The method comprises using at least 2 sets of hybridisation probes, where at least comprises one or more PNA probes capable of hybridisation to chromosome. The methods can be used for the detection of chromosome chromosome. The methods can be used for the diagnosis of disorders and diseases related to chromosomal aberrations or abnormalities such as e.g. cheated to chromosomal aberrations or abnormalities such as e.g. chempositic malignancies, cancers and inborn constituel diseases. The method may be used for detecting viral sequences and their localization in the chromosome. In plant biology, the methods can be used for monitoring the efficiency of transferring herbicide resistance genes to a
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The invention relates to detecting (MI) polymorphisms in a uridine diphosphate glucuronosyltransferase (UGT) gene promoter by determining the presence of five thymidine-adenine (TA) repeats in the promoter, where the presence of the five TA repeats correlates with increased expression of the gene. The method is used for detecting polymorphisms in useful for screening individuals for variation in glucuronidation activity, for optimising drug dosages for a patient, where the drugs (GG. Irinotecan or TAS-10) are glucuronidated by UGT (preferably UGTIA1) and the activity of the drug is effected by its level of glucurodination. The method preferably involves obtaining DNA from an
and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the MGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, comprises determining the presence of five thymidine-adenine repeats in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ds; UGT1A1; promoter; Gilbert's syndrome; hyperbilirubinaemia;
uridine diphosphate glucuronosyltransferase; Crigler-Najjar syndrome;
UGT; polymorphism detection; TA repeat; glucuronidation; Irinotecan;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human UGT1a1 promoter polymorphism (TA)7 repeat region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 5 A; 6 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      at ftp.wipo.int/pub/published_pct_sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1992 CACCTTCAAGCAGCTGG 2008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CACCATCAAGCAGCTGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK90422 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ARCH-) ARCH DEV CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PAS-103; xenobiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rienzo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo gapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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contained in the DNA and determining the number of TA repeats in the promoter. Contained in the DNA and determining the number of TA repeats in the promoter. Thus the DNA being amplified comprises all or part of UGTIAI promoter. The DNA is amplified by a polymerase chain reaction and sequencing the amplified DNA. The polymorphism comprises an allele consisting of five TA repeats (TA) s. six TA repeats (TA) s. or seven TA repeats (TA) s. (TA) 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, comprises determining the presence of five thymidine-adenine repeats in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, ds, UGT1A1, promoter, Gilbert's syndrome, hyperbilirubinaemia, uridine diphosphate glucuronosyltransferase, Crigler-Najjar syndrome, UGT, polymorphism detection, TA repeat, glucuronidation, Irinotecan;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present sequence is the UGT1A1 promoter (TA) 7 repeat region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human UGTlal promoter polymorphism (TA)7 repeat region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 9 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.4; DB 1;
Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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nes 16; Conservative
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cosperssion of the gene. The method is used for detecting polymorphisms in a UGT gene promoter, preferably a UGT I (UGTIAI) gene promoter. (MI) is a UGT gene promoter, preferably a UGT I (UGTIAI) gene promoter. (MI) is activity, for optimising dosages for a patient, where the drugs cativity, for optimising dosages for a patient, where the drugs (e.g. Irinotecan or TAS-103) are glucuronidated by UGT (preferably UGTIAI) and the activity of the drug is effected by its level of glucurodination. The method preferably involves obtaining DNA from an individual, amplifying all or part of a UGT gene promoter (UGTIAI) gene promoter. Thus the DNA being amplified comprises all or part of the promoter. The DNA is amplified comprises all or part of the number of TA repeats is determined by gel electrophoresis or by UGTIAI promoter. The DNA is amplified by gel electrophoresis or by the number of TA repeats is determined by gel electrophoresis or by CC generaling of five TA repeats (TA) 5, six TA repeats (TA) 6, or seven TA repeats (TA) 7. The promoter has any one of the genotypes (TA) 5/(TA) 8. (TA) 7/(TA) 8. (TA)

Sequence 17 BP; 9 A; 0 C; 0 G; 8 T; 0 U; 0 Other;

Gaps ; 0.4%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03; 1; Indels 0; Mismatches 94.18; Local Similarity 94.1 tes 16; Conservative Query Match Matches

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3462 TTATATATATCTATATA 3478 17 rrarararararara

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RESULT 1186 **AAL50679**

AAL50679 standard; DNA; 17 BP AAL50679;

(first entry) 16-JAN-2003 Human uridine diphosphate glucuronosyltransferase gene polymorphism #13.

Human; polymorphism; TA repeat; ds; UGT; thymidine-adenine repeat; uridine diphosphate glucuronosyltransferase gene promoter; UGT1A1; drug dosage optimisation; xenobiotic sensitivity.

Homo sapiens

US2002115097-A1

22-AUG-2002

01-FEB-2002; 2002US-00061693

99US-00251274. 16-FEB-1999;

(ARCH-) ARCH DEV CORP.

Ratain MJ; Rienzo AD, Iyer L,

WPI; 2002-740095/80.

Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, involves determining number of thymidine-adenine repeats in the promoter.

Example 6; Page 2; 13pp; English

The invention comprises a method for detecting polymorphisms in a uridine diphosphate glucuronosyltransferase (UGT) gene promoter (preferably UGT1A1). The method involves determining the number of thymidine-adenine (TA) repeats in the promoter — as the number of TA repeats correlates with expression of the UGT gene. The method of the invention is useful for detecting polymorphisms in a UGT gene promoter. The method of the invention is also useful in optimising drug dosages and predicting an individual's sensitivity to xenobiotics for drugs and xenobiotics that are glucuronidated by UGT. The present DNA sequence represents a UGT gene TA repeat polymorphism \$\$666666666668\$\$

Sequence 17 BP; 9 A; 0 C; 0 G; 8 T; 0 U; 0 Other;

Gaps .; 0 Score 15.4; DB 1; Length 17; Pred. No. 1.2e+03; 0; Mismatches 1; Indels 0.4%; 16; Conservative Query Match Best Local Similarity Matches

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RESULT 1187 AAL50679,

AAL50679 standard; DNA; 17 BP

AAL50679;

(first entry) 16-JAN-2003 Human uridine diphosphate glucuronosyltransferase gene polymorphism #13.

Human, polymorphism; TA repeat, ds, UGT; thymidine-adenine repeat, uridine diphosphate glucuronosyltransferase gene promoter; UGT1A1, drug dosage optimisation; xenobiotic sensitivity.

Homo sapiens.

US2002115097-A1.

22-AUG-2002

01-FEB-2002; 2002US-00061693.

16-FEB-1999;

(ARCH-) ARCH DEV CORP

Ratain MJ Rienzo AD, Iyer L,

WPI; 2002-740095/80.

Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, involves determining number of thymidine-adenine repeats in the promoter.

Example 6; Page 2; 13pp; English.

The invention comprises a method for detecting polymorphisms in a uridine diphosphate glucuronosyltransferase (UGT) gene promoter (preferably UGTLA1). The method involves determining the number of thymidine-adenine (TA) repeats in the promoter - as the number of TA repeats correlates with expression of the UGT gene. The method of the invention is useful for detecting polymorphisms in a UGT gene promoter. The method of the invention is also useful in optimising drug dosages and predicting an individual sensitivity to xenobiotics for drugs and xenobiotics that are glucuronidated by UGT. The present DNA sequence represents a UGT gene TA repeat polymorphism

Sequence 17 BP; 9 A; 0 C; 0 G; 8 T; 0 U; 0 Other;

Query Match

DB 1; Length 17; 0.4%; Score 15.4; ö

Gaps

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Length 17; Indels

Score 15.4; DB 1; Pred. No. 1.2e+03; 0; Mismatches

0.4%;

Query Match 0.4 Best Local Similarity 94.1 Matches 16; Conservative

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Seguence 17 BP; 2 A; 8 C; 6 G; 1 T; 0 U; 0 Other;

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence of 6711, (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner (I) and nucleic acids (II) for identifying a specific binding partner (I) and nucleic acids (II) caused by altered expression of human POSHL1 including diagnosing and treating caused by altered expression of human POSHL1 including diagnosing and treating caused by altered expression of numan POSHL1 including diagnosing and treating caused by altered expression of human post in general partner of vaccines and (II) is treating caused by altered expression of producing the proteins. The transgenic non-human animals capable of producing the proteins. The present sequence did not form part of the privanted specification, but is based on sequence information supplied to berwent by the European Patent Office
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                        Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
                                   Gaps
                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; SEQ ID NO 442; 60pp + Sequence Listing; English.
Pred. No. 1.26+03;
                                                                                                                                                                                                                                                                                                        Human POSHL1 scanning oligonucleotide SEQ ID NO 442.
                                  0; Mismatches
                                                                     3462 TTATATATATCTATATA 3478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US0006654.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US0006667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
10-OCT-2001; 2001US-00806670.
                                                                                                                                                                                                  ABV89729 standard; DNA; 17 BP.
                                                                                                28-JAN-2002; 2002EP-00001165
               94.18;
                                                                                                                                                                                                                                                                        (first entry)
                 Best Local Similarity 94.1
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP1239051-A2.
                                                                                                                                                                                                                                                                        23-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shannon M;
                                                                                                                                                                                                                                    ABV89729;
                                                                                                                                                              1188
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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, liver failure, hepatocaliular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5 terminal nucleocides and a 3 and modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                         WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; lifertion; tancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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0.4%; Score 15.4; DB 1; Length 17;
Best Local Similarity 94.1%; Pred. No. 1.2e+03;
Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                       WNV minus strand Inozyme substrate SEQ ID NO 11304.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 6 A; 5 C; 4 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 23; SEQ ID NO 11304; 495pp; English.
 ВР
                                                                                                                                                                                                                                                                                                                                                                 19-OCT-2001; 2001WO-US048350.
                                                                                                                                                                                                                                                                                                                                                                                                   20-OCT-2000; 2000US-0242411P.
ACN11301 standard; RNA; 17
                                                                     22-APR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM (BLAT/) BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen JA;
                                                                                                                                                                                                                    Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-706994/76.
                                                                                                                                                                                                                                                    West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                    BLATT L.
                                                                                                                                                                                                                                                                                          WO200268637-A2.
                                                                                                                                                                                                                                                                                                                             06-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blatt L,
                                   ACN11301;
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1820 TCCTGCTCTGGGAGATC 1836

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Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.

Tumour suppression related human fukutin oligo SEQ ID No 5813.

(first entry)

13-JUN-2003

ABT40176;

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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WWV). The nucleic acid molecules are useful for treating a condition related to WWV infection e.g. pancreatitis, encephalitis, waycarditis, meningitis, neurologic infection, hepatitis, neurologic infection, hepatitis, molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least the 2'-O-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nucleotides and a 3' and modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 47502-17514 are not given molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                 virucide, neuroprotective, antibacterial, replication, pancreatitis, encephalitis, myocarditis, meningitis; infection, hepatitis, liver failure, cancer, cirrhosis, Hammerhead, Inozyme, DNAzyme,
                                                                                                                                                                                                                                             WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 2 A; 4 C; 5 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 23; SEQ ID NO 2136; 495pp; English.
                                                                                                                                                                                                              WNV Inozyme substrate SEQ ID NO 2136.
17 TTCTGCTCTGGGAGATC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-2001; 2001WO-US048350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-OCT-2000; 2000US-0242411P
                                                                                                       ACN02146 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 64...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatt L, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                     Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-706994/76.
                                                                                                                                                                                                                                                                                                                                                    West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                     WO200268637-A2.
                                                                                                                                                                            22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                         06-SEP-2002.
                                                                                                                                         ACN02146;
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                                                                        RESULT 1190
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New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies

Tuijnder M;

Amson R,

Telerman A,

WPI; 2003-313353/30.

(MOLE-) MOLECULAR ENGINES LAB

17-SEP-2001; 2001FR-00011978. 17-SEP-2002; 2002WO-IB004208.

WO2003025175-A2 Homo sapiens.

27-MAR-2003

Disclosure; Page 713; 720pp; French.

and transfected cells.

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The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that thybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic caids of the invention are useful as probes and primers for detecting, dentifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, containing the nucleic acids, cells containing the cvector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell cegases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in disease whe nother samples is useful for diagnosis and/or prognasis of these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 6 A; 5 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ65192 standard; RNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATCCCCAACAATGTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ65192;
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ABZ65192
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RX H
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Gaps

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Length 17; 1; Indels

0.4%; Score 15.4; DB 1; 64.7%; Pred. No. 1.2e+03; iive 5; Mismatches 1;

1820 TCCTGCTCTGGGAGATC 1836

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: |:||:|:||:| UUCUGCUCUGGGAGAUC

ABT40176 standard; DNA; 17

RESULT 1191 ABT40176 ID ABT40176

vivlemore401-10.rng

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in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the development of compounds with a cytostatic, virucide, neuroprotective, nootropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one component of a gene chip, in vitro as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of planamecuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence data for this patent did not form part of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to novel isolated nucleic acid sequences involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPOUS; Rep-X; human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1; Length 17;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 6 A; 5 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      at ftp.wipo.int/pub/publishedpct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 4181; 30pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2206 GGTCCCCAACAATGTGA 2222
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                                                                                                                                                                                                                                                                                                                                                           (MOLE-) MOLECULAR ENGINES LAB
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                                                                                                                                                                                              17-SEP-2002; 2002WO-IB004523.
                                                                                                                                                                                                                                                                               17-SEP-2001; 2001FR-00011980.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%;
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Best Local Similarity 94.1
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                           relerman A, Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-313354/30.
                                      WO2003025177-A2
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                                                                                                                  27-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel short interfering RNA and enzymatic nucleic acid useful for
                                                                                                                                                                                          Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 1 A; 5 C; 7 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Page 145; 185pp; English.
                                                                                                                  Human HER2 DNAzyme substrate #649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1678 GACTTCGGGCTGGCCCG 1694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADI51678 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAY-2001; 2001US-0294140P
06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAY-2002; 2002WO-US016840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ribozymes of the invention
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                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200297114-A2.
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ното варіепв
                                      21-MAR-2003
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AD151678
XX
AC AD151678
AC AD151678
DT 15-APR-;
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DT 15-APR-;
XX
KW tumour 8
KW cytostaf
KW SYGOSTAF
KW CYTOSTAF
KW CONTROL
KW CONT
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The invention discloses a method for identifying a candidate polymorphic repeat within a coding sequence (expressed sequence tag, BST), which comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, FOMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                              diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's araxis, myotonic dystrophy, hyperandrogenemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Simple sequence repeat, plant, ds, SSR, ryegrass, fescue, tandem repeat, cereal profiling, grass profiling; seed batch purity testing.
                                                                                                                                                          Identifying a candidate polymorphic repeat within a coding sequence, funderstanding or treating genetic disease, comprises detecting tandem repeats in a target coding sequence and scoring the repeats for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.4; DB 1; Length 18; 94.1%; Pred. No. 1.3e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 8 A; 0 C; 1 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE SOUTH AUSTRALIA SOUTH AUSTRALIAN R. UNIV SOUTHERN CROSS.
STATE VICTORIA DEPT NATURAL RES & ENVIRO.
                                                                                         Fondon JW
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INT MAIZE & WHEAT IMPROVEMENT CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Simple sequence repeat, SSR, #30.
                                                                                                                                                                                                                                                    Example; Col 385; 588pp; English.
                                                                                       Garner HR, Wren JD, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2824 ATATATACATATATA 2840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS13733 standard; DNA; 18 BP.
                  99US-00475947.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-DEC-1999; 99AU-00004906.
04-MAY-2000; 2000AU-00007310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94.18;
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                                                  (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                               polymorphic probability.
                                                                                                                        WPI; 2003-208818/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                31-DEC-1999;
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(VICT-)
(UYAD-)
(ITMA-)
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The invention relates to a substantially purified or isolated nucleic acid (I) from ryegrass or fescue species including a simple sequence crepeat (SSR), having 2 or more tandemly repeated nucleotide core elements C2 -6 nucleotides in length. Also included are a nucleic acid primer can table for amplifying an SSR, identifying (MI) an SSR by preparing a library of ryegrass or fescue genomic DNA enriched for SSRs and identifying clones in the library containing SSRs, a library of ryegrass or fescue genomic DNA enriched for SSRs prepared by the MI, selecting for a gene in grass or cereal breeding by identifying an SSR that is closely a sesociated with the gene such that the SSR and the gene are core associated with the gene such that the SSR and the gene are preferentially co-inherited, and selecting for the SSR in the breeding, a method for DNA profiling grass or cereal species varieties by assessing variation within seed batch of an SSR. The SSRs can be used in the selection of genes in grass or cereal breeding, for profiling grass or cereal species varieties, for testing the purity of grass or cereal seed batches, and for DNA profiling to establish the calesing the grass or cereal seed batches, and for DNA profiling to establish the calesing the grass or cereal seed batches, and for DNA profiling to establish the sequence is a ryegrass or fescue SSR
                                                                                New simple sequence repeats having 2 or more tandemly repeated nucleotide core elements isolated from ryegrass and fescue, useful for selecting of genes in grass or cereal breeding or profiling grass or cereal species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Development of polymerase chain reaction-single strand conformation polymorphism(pcr-sscp) to identify korean beef cattle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bovine; melanocortin; receptor; BDF3; PCR; primer; cattle; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.4%; Score 15.4; DB 1; Length 18; Best Local Similarity 94.1%; Pred. No. 1.3e+03; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 9 A; 8 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bovine melanocortin receptor BDF3 PCR primer #2.
                                                                                                                                                                                         Claim 6; Page 51; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL53363 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JUL-2002 (first entry)
  Forster JW, Jones ES;
                                         WPI; 2001-512563/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-495297/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KR2001016520-A.
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                                                                                                                                                varieties
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gene fragments from CACP genomic DNA or to amplify cDNA

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The composition may further comprise a local anaesthetic. The composition of the invention may be administered via intra-articular or intravenous injection. The human CACP protein is identified in the invention as being megakaryocyte stimulating factor (MSP). The gene encoding CACP protein (MSP) is located on chromosome 1q25-31, and mutations in this gene are responsible for the heritable disorder camptodactyly- arthropathy-coxa vara-pericarditis, in which patients have synovial hyperplasia without evidence of inflammation. CACP protein (MSP) acts as a synovium lubricant, and can be used to lubricate tissue and joints in the treatment of osteoarthritis in the composition may be applied to reduce the cymptoms of osteoarthritis (e.g., joint pain, loss of range of movement or joint damage). Sequences AAF59672-AAF59693 represent PCR primers used
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                             The present invention relates to a new technique of polymerase chain reaction-single strand conformation polymorphism (PCR-SSCP) to identify accurately Korean beef cattle in a short time by removing processes which requires expensive restriction enzymes. The present sequence is a PCR primer for bovine melanocortin receptor (BDF3) which was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, CACP protein, camptodactyly-arthropathy-coxa vara-pericarditis, MSF; megakaryocyte stimulating factor; synovial lubricant; chromosome 1q25-31; osteoarthritis; joint lubrication; osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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pericarditis protein in combination with an anesthetic, useful for
treating osteoarthritis, or as lubricants of tissue and joints.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ó
                                                                                                                                                                                                                                                                                                                                                                                  Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 3 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                              Score 15.4; DB 1;
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human CACP (MSF) gene exon 6 reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYCA-) UNIV CASE WESTERN RESERVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2088 CCCGGGTGGCCAGGACA 2104
   Claim 1; Page 10; 10pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTGGGTGGCCAGGACA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF59685 standard; DNA; 18 BP
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                                                                                                                                                                                                                                                                                                                                                                                  0.4%;
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19-JUL-2000; 2000US-00145328
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                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200107068-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF59685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; hypersensitivity disease; infectious disease; neoplastic disease; disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; parasitic infection; schistosome; filaria, bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; T-cell associated disease; Vbeta; autoimmune disease;
                                                                                       0.4%; Score 15.4; DB 1; Length 18; 94.1%; Pred. No. 1.3e+03;
                                                                                                                                1; Indels
                                                      Sequence 18 BP; 2 A; 2 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 983; 164pp; English.
to amplify exonic gene fragments from CA fragments for the detection of mutations
                                                                                                                                                                                                                                                                                                                                                                                                                Human Vbeta gene repeat sequence #579.
                                                                                                                                                                     2600 CCCACACCCAAAGCTGA 2616
                                                                                                                                                                                                                                                                                                     BP.
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95US-00531241.
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                                                                                                                                                                                          18 CCTACACCCAAAGCTGA
                                                                                                                                                                                                                                                                                                     ADH70789 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2004 (first entry)
                                                                                                                                Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-059052/06
                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  breast cancer; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HOOD/) HOOD L E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2002150891-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                          ADH70789;
                                                                                             Query Match
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Gaps

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allergies, Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by viruses such as HIV, fungal infections such as those caused by the yeast genus Candida, parastic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; T-cell associated disease; Wheta, autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease, hypersensitivity disease; type I hypersensitivity; doodpasture's syndrom; type II hypersensitivity; doodpasture's syndrom; type IV hypersensitivity; doodpasture's syndrom; type IV hypersensitivity; alberton disease; viral infection; filaria; bacterial infection; Mycobacterium; neoplastic disease; leukaemia; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                    such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.4; DB 1; Length 18; 14.1%; Pred. No. 1.3e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 9 A; 8 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human Vbeta gene repeat sequence #23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2335 GTGTGTGTGTGTGTG 2351
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95US-00531241,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.4
Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2002150891-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, VbetaRNA or cDNA. The kit is useful for diagnosing organ transplant

Disclosure; SEQ ID NO 427; 164pp; English.

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proluding autoimmune disease, degenerative nervous system diseases, graft versus host disease, hypersensitivity diseases, infectious diseases and neoplastic diseases. Autoimmune diseases include Addison's disease, attrophic gastritis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include Type allergies, Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by whithes such as HIV, fungal infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic infections such as those caused by Mycobacterium, Neoplastic diseases include lymphoproliferative diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 18 claims primers for use in detecting mutations in a mammalian gene for a structural protein of cartilage comprising a sequence identified in Table I (Page 18-31). Table I includes 179 primer sequences (see AAQ65728-Q65906). The following details are given for primer 103: Region/exon: 13 Direction: sense Primer position: 2165 (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diseases
and
                                                                                                                                                                                                                                                                   such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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    diagnosing and treating T-cell associated diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting genetic pre-disposition to osteoarthritis - and other involving mutation in cartilage protein genes, by amplification analysis of DNA and comparison with standards.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Type II procollagen; COL2A1; amplification; primer; polymerase chain reaction; PCR; osteoarthritis; cartilage; ss.
                                                                                                                                                                                                                                                                                                                                                                 Score 15.4; DB 1; Length 19; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
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Hopkinson I, Ahmad NN;
                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 10 A; 0 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Type II procollagen sequencing primer 103.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            2836 ATATATATACATATAT 2852
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                                                                                                                                                                                                                                                                                                                                                                                   11 Similarity 94.1%;
16; Conservative
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(first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-1992;
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19-DEC-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tomato, TDET1 (HP-2) gene; light hypersensitive phenotype; mutation; carotenoid; chlorophyll; flavonoid; transgenic; TDET1 gene; anthocyanin; agro-industrial; antioxidant; antitumoural; plant protection; variant; ornamental; herbicide; fruit ripening; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequences of the tomato TDET1 (HP-2) gene, which if modified,
                                                                         Gaps
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0.4%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.46+03; ive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tomato TDET1 gene amplifying nested primer.
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                                                                                                                                                                                                                                                                                                                                                                                                        AAX60990 standard; cDNA to mRNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 15; 57pp; English
                                                                                                                                              2991
                                                                                                                                                                                            AGATGACCAGGGCTTTT 18
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                                                                                                                                              2975 AGAGGACCAGGGCTTTT
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Query Match
Best Local Similarity 94.1
Matches 16; Conservative
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The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of the SCD (stearoyl-CoA desaturase) gene genes in interference. Also described: (1) modulating expression of SCD genes in cells, tissue explants or organisms by introduction of siNA: kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA. SCD inhibiting siNAs have ancrectic, antidiabetic, antiatreriosclerotic, cytostatic and virucide activities. The siNAs can be used to modulate expression of SCD genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target studying gene function and validation; genetic empinesting; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which is
                                                                                                                                                                                                                                                                  short interfering nucleic acid; siNA; downregulation; inhibition; SCD; stearcyl-CoA desaturase; RNA interference; anorectic; antidiabetic; antiatreiosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid, useful e.g. for treatment and diagnosis of obesity or diabetes, downregulates expression of the stearoyl-CoA desaturase gene.
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Pred. No. 1.4e+03;
7; Mismatches 1; Indels
                                                                                                                                                                                                                                  Stearoyl-CoA desaturase siNA oligonucleotide SEQ ID NO:134.
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                                                                                                                          BP
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09-SEP-2002; 2002US-0409293P.
20-SEP-2002; 2002US-0412304P.
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2002US-0386782P.
2002US-0406784P.
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Similarity 52.9%;
9; Conservative
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1291 GCCGTGAAGATGCTGAA
                                   19 GCCGTGAAGATGATGAA
                                                                                                                        ADE27190 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                29-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-721687/68.
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                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003070885-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                            ADE27190;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                        RESULT 1202
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Matches
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Gaps

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0.4%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.4e+03; tive 0; Mismatches 1; Indels

Query Match
Best Local Similarity 94.1
Matches 16; Conservative

19 TATATACATATACAT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of the SCD (stearoyl-CoA desaturase) gene by RNA interference. Also described: (1) modulating expression of SCD genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA. SCD inhibiting siNAs have anorectic, antidiabetic, antidiates and antidiate expression of SCD inhibiting virucide activities. The siNAs can be used to modulate expression of SCD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 genes, in cells, tissue explants or organisms, e.g. for treating obesity diabeters (types I and II); atherosolerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD sina, which is used in the exemplification of the present invention.
                                                                                                                                                                                          short interfering nucleic acid; siNA; downregulation; inhibition; SCD; stearcyl-CoA desaturase; RNA interference; anorectic; antidiabetic; antiarteriosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for treatment
expression of
                                                                                                                                                                   Stearoyl-CoA desaturase siNA oligonucleotide SEQ ID NO:424.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 8 A; 1 C; 2 G; 0 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New short interfering nucleic acid, useful e.g. diagnosis of obesity or diabetes, downregulates stearoyl-CoA desaturase gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; SEQ ID NO 424; 139pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J, Beigelman L, Thompson J;
2825 TATATACATATATAT 2841
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                                                                                                                                                                                                                                                                                                                                                                                                              2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-SEP-2002; 2002US-0412304P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                13-FEB-2003; 2003WO-US004317
                                                                                                                                                                                                                                                                                                                                                                          2002US-0358580P
                                                                                       ADE27480 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-721687/68.
                                                                                                                                                                                                                                                                                              WO2003070885-A2.
                                                                                                                                                                                                                                                                                                                                                                                                              29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                      11-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                         09-SEP-2002;
                                                                                                                                          29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                          20-FEB-2002;
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                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                ADE27480;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      당
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene function studies and gene mapping. The current sequence is that the human BCR-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                               short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; BCR; ss; siRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ô
                                                                                                                                                                                                                            Human breakpoint cluster region-targeted siRNA - SEQ ID 410.
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4.1%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 11 C; 1 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 7; SEQ ID NO 410; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chowrira B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-FEB-2002; 2002US-0358580P.
11-WAR-2002; 2002US-0361124P.
06-JUN-2002; 2002US-0386782P.
15-AUG-2002; 2002US-0406784P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-04083P.
14-JJN-2003; 2003US-0439323P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 855 GGAGGAGCTGGTGGAGG 871
                                                        BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF83853 standard; RNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.1%;
                                                        ADF84116 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17 GGTGGAGCTGGTGGAGG
                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-679889/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003070972-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen J,
                                                                                                                                                                       26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-AUG-2003
                                                                                                              ADF84116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
RESULT 1204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1205
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ID ADF8
XX
                             ADF84116
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Query Match
Best Local Similarity 94.1
Matches 16; Conservative

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0; Gaps

1; Indels

0; Mismatches

0.4%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.4e+03;

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The invention describes the use of a nucleic acid (I), of at least 8 nucleotides (nt), and (essentially) identical with nucleic acid (II) present at specified positions in chromosomes 5 or 11 of pigs (or homologous positions in other animals) for determining predisposition for expression or inheritance of the 'splay-leg' phenotype in mammals and poultry. The specified positions are regions of microsatellite markers (a) for chromosome 5, 84426, 8425, 84995, IGF1, 80005, 84933, 841987, 80108, 84325, 84995, IGF1, 80005, 84983, 841987, 80108, 8435, 801094, 8496, Also described are: an in vitro method for determining the predisposition for expression or inheritance of the 'splay-leg' phenotype in mammals, their (un)fertilized eggs or sperm by detectining presence, condition or expression of (II); and a kit comprising primer pair for amplification of (II); and a kit comprising primer pair for amplification of (II); in one or more containers. The method is used for actual binds to (II), in one or more containers. The method is used for eatle, dogs, pigs, chickens etc.) that lack predisposition to the phenotype, particularly as a genomic screen applied to many mammals in a population. This sequence represents a primer that can be used to be account of the 'splay-leg' determine predisposition for expression or inheritance of the 'splay-leg'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       splay-leg phenotype; microsatellite marker; Sw1468; Sw2; Sw1200; Sw2425; Sw995; IGP1; S0005; Sw963; Sw1987; S0018; Sw904; Sw310; Swr1974; Swr426; Sw1094; Sw986; Sw1982; S0182; S0071; Sw2008; Sw435; S0009; S0230; Sw486; selective breeding; ss; primer.
Sw1094; Sw986; Sw1982; S0182; S0071; Sw2008; Sw435; S0009; S0230; Sw486;
                                                                                                                                                                                                                                                                                                                                                    Determining predisposition to the splay-leg phenotype in mammals and birds, useful for selecting breeding animals, by detecting specific chromosomal markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Splay-leg phenotype inheritance pattern associated primer seqid 85.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.4e+03;
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                                                                                                                                                                                                                                         (FBFF-) FBF FOERDERVEREIN BIOTECHNOLOGIEFORSCHUN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 2 A; 3 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 36; 104pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3212 CCCTCCAAGCCTAAAAG 3228
                   selective breeding; ss; primer.
                                                                                                                                                                                                                                                                              Schwarz S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                 10-OCT-2003; 2003WO-EP011269.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       m
                                                                                                                                                                                                      11-OCT-2002; 2002EP-00022810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 94.1%;
nes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                              Kalm E, Reinsch N,
                                                                                                                                                                                                                                                                                                                 WPI; 2004-348471/32
                                                                                         WO2004035820-A1
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                                                                                                                              29-APR-2004
                                                      Sus scrofa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADN08143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-vabl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siNNA of the invention demonstrates cycostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human BCR-targeted siNNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         splay-leg phenotype; microsatellite marker; Sw1468; Sw2; Sw1200; Sw2425;
Sw995; IGF1; S0005; Sw963; Sw1987; S0018; Sw904; Sw310; Swr1974; Swr426;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
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                                                                                                       short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; BCR; ss; siRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Splay-leg phenotype inheritance pattern associated primer seqid 36
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                                                                       Human breakpoint cluster region-targeted siRNA - SEQ ID 147.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chowrira
                                                                                                                                                                                                                                                                                                                                 .20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0353124P.
06-JUN-2002; 2002US-0386782P.
15-AUG-2002; 2002US-0404039P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-040923P.
14-JAN-2003; 2003US-043922P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 GGUGGAGCUGGUGGAGG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADN08094 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                               20-FEB-2003; 2003WO-US005234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-2003; 2003US-0440129P
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Best Local Similarity 82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-679889/64.
                                                                                                                                                                                                                       WO2003070972-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUL-2004
                                                                                                                                                                                    Homo sapiens
                                    26-FEB-2004
                                                                                                                                                                                                                                                            28-AUG-2003
ADF83853;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADN08094;
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Gaps

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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding fibroblast growth factor (RGP) receptor 3 (also known as FGR-3) ACH, JTK4 and CEK2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with FGRR-3 such as developmental disorders or hyperporliferative disorders, especially cancer of colorectal, bladder, bone, lung, cervical, brast or skin. They are useful as research reagents, therapeutics, prophylaxis, kits and diagnostics, and as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion of the genes expressed within cells and tissues. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human FGFR-3
                                                                    /*tag= a //mod base= OTHER //mod base= OTHER //mod base= OTHER //mode are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel compound targeted to a nucleic acid molecule encoding fibroblast growth factor receptor 3, useful for inhibiting the expression of the receptor and for treating an animal having cancer or developmental
                                                                                                                                                             1. .5
/*tag= b
/mod_base= OTHER
16. .20
                                                                                                                                                                                                                                                                                                                                        /note= "2 -methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.4; DB 1; Length 20; Pred. No. 1.4e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 9 A; 2 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; Page 79; 120pp; English.
                    location/Qualifiers
                                                                                                                                                                                                                                                                                                                   /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3464 ATATATATATATA 3480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ATATATGTATATATA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABT07496 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-SEP-2002; 2002WO-US028549.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .0-SEP-2001; 2001US-00953047.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                       /*tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 94.1
Matches 16, Conservative
                                                               /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-313244/30.
                                                                                                                                                                                                                                                                                                                                                                                           WO2003023004-A2
                      Key
modified base
                                                                                                                                                                     modified base
                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1209
8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes the use of a nucleic acid (I), of at least 8

nucleotides (nt), and (essentially) identical with nucleic acid (II)

present at specified positions in chromosomes 5 or 11 of pigs (or
homologous positions in other animals) for determining predisposition for
expression or inheritance of the 'splay-leg' phenotype in mammals and
poultry. The specified positions are regions of microsacellite markers
(a) for chromosome 5, 8w168, 8w2, 8w1200, 8w425, 8w995, IGT, 80005,
8w963, Sw1987, S0018, Sw904, Sw110, Sw1200, Sw425, Sw995, IGT, 80009,
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Sw1987, S0018, Sw904, Sw110, Sw1974, Swr426, Sw1094, Sw986 or
Sw963, Diddention or expression or inheritance. The method is used
for selection of animals (domestic, breeding or farm animals, e.g.
cattle, dogs, pigs, chickens etc.) that lack predisposition to the
population. This sequence represents a primer that can be used to
cheternine predisposition for expression or inheritance of the 'splay-leg'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; antisense; fibroblast growth factor receptor 3; prophylaxis; developmental disorder; hyperproliferative disorder; antisense therapy; FGFR-3; ACH; JTK4; CEK2; cancer; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                             Determining predisposition to the splay-leg phenotype in mammals and birds, useful for selecting breeding animals, by detecting specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 15.4; DB 1; Length 19;
4.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human FGFR-3 antisense oligonucleotide, ISIS #125205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 2 A; 3 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                              (FBFF-) FBF FOERDERVEREIN BIOTECHNOLOGIEFORSCHUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 62; 104pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3212 CCCTCCAAGCCTAAAAG 3228
                                                                                                                                                                                                                                                                                                              Schwarz S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD55499 standard; DNA; 20 BP.
                                                                                                                                                                10-OCT-2003; 2003WO-EP011269.
                                                                                                                                                                                                                11-OCT-2002; 2002EP-00022810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19 cccrccaacccrcaaag 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 94.1%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-AUG-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               chromosomal markers.
                                                                                                                                                                                                                                                                                                              Reinsch N,
                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-348471/32
                                                                  WO2004035820-A1.
                                                                                                               29-APR-2004
                    Sus scrofa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD55499;
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Gaps ö Cytostatic, antidiabetic; antisense therapy; aberrant insulin regulation; protein phosphatase 2 catalytic beta subunit; antisense compound; cancer;

Homo sapiens. Synthetic.

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Rat protein phosphatase 2 oligo inhibitor SEQ ID No 110.

Homo sapiens.

dß.

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The invention relates to a novel compound 8-50 nuclectides in length targeted to a nucleic acid molecule encoding a protein phosphatase 2 catalytic beta subunit, where the compound specifically hybridises with and inhibits the expression of protein phosphatase 2 catalytic beta subunits, or specifically hybridises with at least an 8-nucleotide subunits, or specifically hybridises with at least an 8-nucleotide consorted or a nucleic acid molecule encoding a protein phosphatase 2 catalytic beta subunit. The antisense compounds are useful for modulating the expression of protein phosphatase 2 catalytic beta subunits and for treating diseases or conditions associated with expression of protein phosphatase 2 catalytic beta subunits, e.g. or parent insulin regulation or diabetes or a hyperproliferative disorder, particularly cancer. The antisense compounds are also useful for diagnostics, therapoutics, prophylaxis, e.g. to prevent or delay inflammation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. This polynuclectide sequence represents an oligonucleotide inhibitor of tat protein phosphatase 2 catalytic beta subunit mRNA levels of the invention. NOTE: This oligonucleotide contains phosphorothioate residues and has 2'- MOE wings with a deoxy gap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides targeted to nucleic acid encoding Protein Phosphatase 2 catalytic subunit beta, useful for treating diseases related to Protein Phosphatase 2 catalytic subunit beta expression, such
hyperproliferative disorder; diabetes; inflammation; tumour; rat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 16; Page 98; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                             31-JAN-2002; 2002WO-US002805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-FEB-2001; 2001US-00780045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia BP, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-657588/70.
                                                                                                    Rattus norvegicus.
                                                                                                                                                                                              WO200264737-A2.
                                                                                                                                                                                                                                                                                              22-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
     SON COURSE OF CO
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Gaps ö 0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03; tive 0; Mismatches 1; Indels Sequence 20 BP; 7 A; 0 C; 1 G; 12 T; 0 U; 0 Other; 3465 TATATATCTATATAT 3481 Best Local Similarity 94.1 Matches 16; Conservative ઠ

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11 1 TATATATGTATATAT

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Human oligonucleotide sequence. ABZ91730 standard; DNA; 20 BP. 17-OCT-2003 (first entry) ABZ91730; RESULT 1210 ABZ91730

Human, antisense; lung dysfunction, nasal airway dysfunction, antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiathmatic; properative; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5, or 3, end genomic flanking regions, 5, and 3, intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of antiinflammatory steroid in a subject, for reducing levels of antiinflammatory steroid in a subject, for reducing levels of cepetor, producing bronchodilation, increasing levels of darnone or lung surfactant in a subject, stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Under The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                              Aguilar D;
                                                                                                                                                                                                                                                                              Katz E, Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 8 A; 0 C; 2 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 6972; 872pp; English.
                                                                                                                                                                                                                                                                              Nyce JW, Li Y, Sandrasagra A, Ka
Miller S, Tang L, Shahabuddin S;
                                                                                                                                        23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                      24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                               (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229219/22.
                                           WO200285308-A2
                                                                                          31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ubiquinone
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Gaps ö Score 15.4; DB 1; Length 20; Pred. No. 1.4e+03; 0; Mismatches 1; Indels AA497002-derived oligonucleotide SEQ ID 6972. 2823 TATATACATATAT 2839 BP. 0.4%; 17 TATATACATACATAT ABD27960 standard; DNA; 20 29-JUL-2004 (first entry) 16; Conservative Query Match Best Local Similarity ABD27960; RESULT 1211 Matches ABD27960/ à

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respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory diserses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

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This probe, from construct M13-mp18, is from 162-181 base of the sequence represented in AAQ05931, which is the presumptive hemolysin gene of L. monocytogenes. The probe is used for detection and enumeration of L. monocytogenes, esp. for detection in dairy prods. such as milk and cheese by colony hybridisation assays. See also AAQ05898-900 and AAQ05930-31. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with orgoing US granted patent numbers. For further information please visit the
                                                                                                                                                                  Listeria monocytogenes; probe ADO8; dairy products; milk; cheese; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic Listeria monocytogenes oligo-nucleotide probes - used for detection and enumeration of organism in dairy prods. e.g. milk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Derwent web site at www.derwent.com/dwpi/updates/ntis_us.html.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 3 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                             Probe AD08 to detect Listeria monocytogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 18; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  446 GCAACTACACCTGCGTC 462
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                                                                                                                                                                                                                                                                                                                                                                               89US-00411965.
                                                                                                                                                                                                                                                                                                                                                                                                                                    89US-00411965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94.1%;
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                              (revised)
(first entry)
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ID AAQ46062 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1990-290094/38.
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                                                                                                                                                                                                                                                                          USN7411965-N.
                                                                                                                                                                                                                                                                                                                                                                                                                                    25-SEP-1989;
                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-1989;
                              17-DEC-2001
16-JAN-1991
                                                                                                                                                                                                                                                                                                                            21-AUG-1990.
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08-FEB-1994
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                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ46062;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Datta A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cheese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1213
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiashmatic, an algesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or maligant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, allergies and/or bronchoconstriction and a sesociated inflammation, allergies and/or surfactant hypoproduction are associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thy midines present in the target RNA serves to prevent the braskdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                         Li Y, Sandrasagra A, .Katz E, Pabalan J, Aguilar D;
Tang L, Shahabuddin S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 8 A; 0 C; 2 G; 10 T; 0 U; 0 Other;
pulmonary transplantation rejection; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 6972; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2823 TATATACATATAT 2839
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                                                                                                                                                                                                               23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                    24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                       (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 94.1%;
ses 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3900/c
AAQ05900 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TATATACATACATAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-093058/08
                                                                                                       WO200285309-A2.
                                                     Homo sapiens
                                                                                                                                                          31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        prevent any
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Miller S,
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Virulence factor; Listeria detection; food poisoning; iap; PCR;
0.4%; Score 15.4; DB 1; Length 20;
94.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90CH-00002190.
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RESULT 1212

AAQ05900

AAQ05900/c ID AAQ09 XX AC AAQ09

Query Match

Matches

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Query Match
Best Local Similarity
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                                                                                                                                                                        Oligos LOI, LO2, LO3 and LO4 are used for the amplification of hly (alpha -haemolysin) virulence factor; and oligos AD07, AD08 and AD09 are used for the amplification of iap (beta-haemolysin) virulence factor. They are used in a detection method for Listeria monocytogenes in food samples which is faster and more sensitive than the classical bacteriological methods. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                Listeria monocytogenes detection by enzymatic nucleic acid amplification - using oligo-nucleotide(s) derived from alpha-haemolysin and/or beta-haemolysin virulence factors in polymerase chain reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Quantitative PCR analysis was performed using primers (AAT31580-81) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Preventing or promoting programmed cell death in vertebrate cells -comprises inhibiting or increasing the activity of interleukin-1-beta converting enzyme, or altering expression of other related genes.
                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ICh-1; ICE-ced-3 homologue; programmed cell death; apoptosis;
interleukin-1 beta converting enzyme; gene therapy; primer; PCR;
polymerase chain reaction; 88.
                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                  0.4%; Score 15.4; DB 1; Length 20;
14.1%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 3 C; 9 G; 6 T; 0 U; 0 Other;

    PCR primer for murine Ich-1 amplification.

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                                                                             Hoefelein C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Page 77; 127pp; English
                                                                                                                                                                                                                                                                                                          446 GCAACTACACCTGCGTC 462
                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                         Claim 3; Page 2; 2pp; German.
         90CH-00002190
                                                                                                                                                                                                                                                                                                                            17 dcaacracaccrecece 1
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                                                                                                                                                                                                                                                                           94.18;
                                                                                                                                                                                                                                                                                                                                                                             AAT31581 Standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                            Best Local Similarity 94.1
Matches 16, Conservative
                                                                            Candrian U, Furrer B,
                                     FURRER B.
HOEFELEIN C.
                            CANDRIAN U.
                                                                                              WPI; 1993-265174/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-333763/33
                                                         LUETHY J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Miura M;
         28-JUN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9620721-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                 AAT31581;
                                    (FURR/)
(HOEF/)
                                                                                                                                                                                                                                                                  Query Match
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                            CAND/)
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amplify mouse Ich-1, primers (AAT31582-83) to amplify human Ich-1, primers (AAT31584-85) to amplify human interleukin-1 beta converting enzyme (HICE), and control primers (AAT31586-87) for mouse beta-actin. Ich-1 is a new member of the ICE/ced-3 family of cell death genes. Ich-11 (See also AAT31552) and Ich-15 (AAT31553) were amplified simulateneously to produce DNA fragments of 234 and 295 bp, respectively. Expression of Ich-18 was detected in HeLa and Jurkat cells but not in THP I or U937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                        cells. Expression levels of Ich-1L increased in dying hybridoma DO11.10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; compound; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ch 0.4%; Score 15.4; DB 1; Length 20; 1 Similarity 94.1%; Pred. No. 1.4e+03; 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Compound simple sequence repeat primer (CA)7.5(TA)2.5.
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                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 8 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genetic trait marking and germplasm comparisons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 84; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ႘
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1353 GGAGATGATGAAGATGA 1369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT30411 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9617082-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03;

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vivlemore401-10.rng

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This 20-mer oligonucleotide primer is used in a novel method of amplifying a nucleic acid molecule involving ribonucleotide tailing of single stranded DNA in the presence of terminal transferase. This method can be used for sequencing RNA, especially the 5'-end and involves a 3' tail of 2 to 4 ribonucleotides being attached onto the target sequence with high efficiency. The first step in cloning and sequencing the unknown 5'-end of a specific mRNA molecule involves the synthesis of single stranded cDNA using a primer specific for a known region in the 3' region of the mRNA. A tail of two rGTPP's is attached to the mRNA using transferase and a specific double stranded DNA adapter is attached via a complementary 3' overhang (i.e. CC) in the presence of T4 crainal transferase and a specific double stranded DNA adapter specific primer and a complementary 3' overhang (i.e. CC) in the presence of T4 crain and a complementary as including a desired restriction site produces a unique PCR product suitable for direct solid-phase stranded of the control of the produces a unique PCR product suitable for direct solid-phase stranded of the control of the produces a unique PCR product suitable for direct solid-phase stranded of the produces a unique PCR product suitable for direct solid-phase stranded of the produces a unique PCR product suitable for direct solid-phase
       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.4; DB 1; Length 20;
Pred. No. 1.4e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ribonucleotide tailing; sequencing; amplification; primer; terminal transferase; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide 20-mer primer for ribonucleotide tailing.
   1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 8 A; 0 C; 3 G; 8 T; 0 U; 1 Other;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sites on the adapter or on the cDNA primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Col 11; 12pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96DE-01001385
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                                                                                                                                                                                                                                                                                                                            AAT89883 standard; DNA; 20
                                                                        2316 TCTGTGTGTGTGTGTGT
                                                                                                                                             17 rargigicierererer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 TTTATATATACATATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 94.1
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MUEL/) MUELLER M W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-365083/34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUL-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                         AAT89883;
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AAX84670
ID AAX8467
                                                                                                                                                                                                                                                           RESULT 1216
   Matches
                                                                                                                                                                                                                                                                                      AAT89883/
XX AAT8
XX AAT8
XX AAT8
XX AAT8
XX KW Ribc
XX Synt
XX Synt
XX Synt
XX COS
XX WPI;
XX WPI;
XX WPI;
XX CC Thi;
CC C Sin;
CC Sin;
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                                                                                        KDR signal transduction inducer protein; human; diabetic retinopathy; vascular endothelial cell growth receptor; abnormal neovascularisation; kinase insert domain-containing receptor; solid tumour proliferation; gene therapy; metastasis; chronic rheumatoid arthritis; psoriasis; retinopathy of prematurity; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense, human Integrin beta 4 binding protein; hIbeta4BP; cytostatic; cell proliferation; cancer; gene therapy; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                            Primer for KDR signal transduction inducer protein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prematurity). It may be used as a screen for candidate KDR signal transduction inhibitors for therapeutic use
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KDR signal transduction inducing protein and antibodies to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.46+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human hIbeta4BP antisense oligonucleotide, ISIS #129429.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 7 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
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1..20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1609 AAGTGCATCCACAGGGA 1625
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                                                                                                                                                                                                                                                                                                    98WO-JP005612.
                                                                                                                                                                                                                                                                                                                                 97JP-00343474.
                                                                                                                                                                                                                                                                                                                                                                HAKKO KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAGTGTATCCACAGGGA
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- AAD35727 standard; DNA; 20
                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-405033/34.
                                                                                                                                                                                                                                                                                                                                                               (KYOW ) KYOWA HAKK(
(SHIB/) SHIBUYA M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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modified_base
                                                                                                                                                                                                      Homo sapiens.
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                               20-SEP-1999
                                                                                                                                                                                                                                                                    24-JUN-1999.
                                                                                                                                                                                         Synthetic.
AAX84670;
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Matches
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The present invention relates to a method for detecting early stage allergic diseases, particularly atopic dermatitis. The method comprises determining the expression levels of granulocyte macrophage colony stimulating factor receptor alpha or beta (GW-CSF R-alpha or -beta), interleukin 3 receptor alpha or beta (GW-CSF R-alpha or -beta). Literleukin 3 receptor Appa (IL-3 R-alpha), bc1-2, bc1-x, platelet activation factor receptor (PAF R) or CD44 in eosinocytes of a subject to be tested. The method further comprises comparison with expression levels in healthy volunteers. The method is useful for the early diagnosis and treatment of early stage allergic diseases such as atopic dermatitis. The present sequence represents a PCR primer used in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting early stage allergic diseases with markers of 7 genes of GM-CSF R-beta, GM-CSF R-alpha, IL-3 R-alpha, PAF R, bcl-2, bcl-x and CD44 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Adipose protein; ss; adp; obesity; transgenic animal; obesity; adipositas; bulimia; wasting; cachexia; eating disorder; obody weight disorder; weight loss; cancer; infectious disease; hypogonadism; prader-Willi syndrome; Laurence-Moon-Biedl syndrome; hypothyroidism; diabetes; Cushing's syndrome; endocrine disorder; protectinal diseases; inflammatory bowel disease; PCR primer; ulcerative colitis; anorexia nervosa; glycogen storage disease; lipid storage disease; lipoma; liposarcoma; heart disease; hypertension; infertility; acquired immunodeficiency syndrome; AIDS.
                                                                              Human, detection of early stage allergic disease, atopic dermatitis, antiallergic, eosinocyte, eosinophil; bcl-2; PCR; primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                       PCR primer #2 for DNA encoding human bcl-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse adipose protein, adp, PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                            (GENO-) GENOX SOYAKU KENKYUSHO KK.
(KOKU-) KOKURITSU SHONI BYOIN INCHO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 19; 25pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1874 TGGAGGAGCTCTTCAAG 1890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                                                                                                                                                                                                                     11-OCT-2000; 2000JP-00311193.
                                                                                                                                                                                                                                                                           11-OCT-2000; 2000JP-00311193
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Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK16356 standard; DNA; 20
(first entry)
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hes 16; Conserv
                                                                                                                                                                                        JP2002119281-A.
                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R-beta, GM-C
eosinophils.
14-AUG-2002
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound targeted to a region of a nucleic acid encoding human Integrin beta 4 binding protein and that inhibits expression of the nucleic acid, for treating e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to antisense compounds targetted to a nucleic acid encoding human. Integrin beta 4 binding protein (Nibeta4BP), which specifically hybridises with the nucleic acid and inhibits its expression. The antisense compounds are useful to prevent or treat diseases associated with Nibeta4BP expression, particularly conditions tinvolving aberrant or deregulated call proliferation (e.g. cancer). The hibeta4BP polynucleotide is used in gene therapy. The present sequence is an antisense oligonucleotide targetted to hibeta4BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "2'methoxyethyl nucleotides"
                                                                                                                          note = "2'methoxyethyl nucleotides
                 /mod_base= OTHER
/note= "Phosphorothioate backbone"
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base= OTHER
                                                                                                       base= OTHER
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/mod_base= m5c
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16. .20
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mod_base= m5c
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Gaps

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Mus musculus

ABK52486 standard; DNA; 20 BP.

ABK52486

RESULT 1219
ABK52486
ID ABK52486
XX
AC ABK52486

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WO200230974-A2 Homo sapiens.

18-APR-2002

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The invention relates to a nucleic acid encoding a adipose (ADP)

polypeptide which regulates, causes or contributes to obesity in an
animal or a human. The polyprucleotides, proteins, ant-adp antibodies,
modulators of adp activity, adp antisense nucleic acids, expression
vectors, adp transgenic animals are useful in the diagnosis and treatment
of obesity, adiposites, bulimia, wasting (cachexia), eating disorders
and/or disorders of body weight/body mass, weight loss due to cancer or
infectious diseases, genetic disorders associated with hypogonadism e.g.
Prader-Willi syndrome, Laurence-Moon-Biedl syndrome, hypothyroidism,
c diabetes, Cushing's syndrome, endocrine disorders, gastrointestinal
diseases, inflammatory bowel disease, ulcerative colitis, and anorexia
nervosa. They are also useful for treating disorders of body weight/mass
c e.g. glycogen storage diseases, and lipid storage diseases and for
treating lipomas, and/or liposarcomas. The compositions are also useful
for treating conditions associated with under weight e.g. enhancing or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             controlling fertility, controlling weight loss in acquired immunodeficiency syndrome (AIDS) or cancer patients. The present sequence is a PCR primer used to amplify an adp nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding adipose polypeptide which regulates, cause or contributes to obesity, useful for treating obesity, heart disease, hypertension, infertility, and controlling weight loss in cancer patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dohrmann C, Haeder T, Rothe M;
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28-JUN-2000; 2000US-0214518P.
17-APR-2001; 2001EP-00109537.
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                                                                                                                                                                                      13-JUN-2001; 2001WO-EP006713
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    WO200196371-A2
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0.4%; Score 15.4; DB 1; Length 20;
44.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
                                                               254 ACAAGAAGCTGCTGGCC 270
                 94.18;
                Best Local Similarity 94.1 Matches 16; Conservative
    Query Match
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Human; NOVX; cytostatic; antiarteriosclerotic; cardiovascular; lymphoma; antidiabetic; immunosuppressive; neuroprotective; gene therapy; cancer; cardiomyopathy; atherosclerosis; cell signal processing; diabetes; AIDS; metabolic pathway modulation; neoplastic; neurological disorder; asthma; adenocarcinoma; prostate cancer; uterus cancer; immune response; crohn's disease; multiple sclerosis; Graft versus host disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Human NOV7 forward PCR primer SEQ ID NO:72.
4 ACAAGAAGCTGCTGTCC 20
                                                                                                                                                                                                   ABN86953 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                             29-JUL-2002 (first entry)
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RESULT 1222 ABT21492/c

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The present invention describes novel human proteins designated NOVX (where X is 1, 2a, 2b, 2c, 2d, 3, 4, 5, 6a, 6b, 7, 8, or 9). NOV1 is a tyrosine kinese felike protein; NOV2a dare keratin 4-like proteins; NOV3 is a collagen-like protein; NOV4 is a cystatin B-like protein; NOV5 is a serotonin receptor-like protein; NOV6 and NOV55 v are cold inducible glycoprotein 30-like protein; NOV7 is a matrilin-like protein; NOV9 is a leukocyte surface antigen (CD53)-like protein; and NOV9 is a tyrosine kinase-like protein. NOV8 sequences have cytostatic, antiatteriosclerotic, cardiovascular; antidiabetic, immunosuppressive and neuroprotective activities, and can be used in gene therapy. The NOVX sequences can be used in therapeutics, particularly for treating, preventing or alleviating a NOVX-associated disorder include state in a subject, particularly a human. These disorders include cratdomyopathy, atherosclerosis, a disorder related to cell signal processing and metabolic pathway modulation or diabetes. The NOVX sequences are also useful for determining the presence of or predisposition to a disease associated with altered levels of NOVX sequences are nucleic acid, particularly cancer. The NOVX sequences are especially useful in therapeutic or prophylactic applications for neuropositic acid, particularly cancer. The NOVX sequences are especially useful in therapeutic or prophylactic applications for neuroposition to a disease associated with altered levels of NOVX respective respecially useful in therapeutic or prophylactic applications for neuropositions in the treatment of include neuropositions and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alsobrook JP, Lepley DM, Burgess CE, Mishra V;
Li L, Padigaru M, Shimkets RA, Zerhusen BD, Spytek KA;
Gerlach V, Macdougall J, Stone D, Gunther E, Ellerman K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             versus host disease. The present sequence represents a PCR primer for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New NOVX polypeptides and polynucleotides, useful for treating or preventing a NOVX-associated disorder or a pathological state in a subject, particularly a human, e.g. cardiomyopathy, atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human NOV7, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            adenocarcinoma, lymphoma, prostate cancer, uterus cancer, immune
response, AIDS, asthma, Crohn's disease, multiple sclerosis or G
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Pred. No. 1.4e+03;
); Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 205; 227pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                                                                                                                                                                                    16-OCT-2000; 2000US-0240648P.
16-OCT-2000; 2000US-0240662P.
16-OCT-2000; 2000US-0240669P.
16-OCT-2000; 2000US-0240703P.
                                                                                                                                                      12-OCT-2001; 2001WO-US031922.
                                                                                                                                                                                                             12-OCT-2000; 2000US-0240113P.
                                                                                                                                                                                                                                      16-OCT-2000; 2000US-0240625P.
                                                                                                                                                                                                                                                                                                                                                                                               16-OCT-2000; 2000US-0240732P.
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16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-444172/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer or diabetes.
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Matches 16; Conserv
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2001US - 0327917P 2001US - 0328029P 2001US - 0328049P 2001US - 0328414P 2001US - 0330414P 2001US - 03310142P 2001US - 034162P 2001US - 034552P 2001US - 034557P 2001US - 034557P

2002US-00262445

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02-OCT-2002; 2002WO-US031357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-) CURAGEN CORP.
                                                                                                                                                                                                              09-OCT-2001; 2
12-OCT-2001; 2
15-OCT-2001; 2
17-OCT-2001; 2
22-OCT-2001; 2
                                                                                                                                                                                                                                                                                                                                                                            29-OCT-2001;
01-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-OCT-2002;
                                                                                                                                                              09-OCT-2001;
09-OCT-2001;
                         17-APR-2003
  The invention relates to a novel method for determining racing potential of a horse. The method comprises measuring: whether grandpaternal DNA is over-represented in the genome of the horse; or in the case where one of the grandmothers was selected for breeding on the basis of racing performance, whether grandmaternal DNA from the selected grandmother is over represented in the genome of the horse which indicates that the horse has good racing potential. The method of the invention is useful for determining the racing potential of a horse or for obtaining a progeny horse with good racing potential. This polynucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Determining the racing potential of a horse comprises measuring whether grandpaternal or grandmaternal DNA from the selected grandmother DNA is over-represented in the genome of the horse.
                                                                                                                                                                                                                Racing potential; horse; grandpaternal DNA; over-represented; breeding; grandmother; performance; progeny horse; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               represents a PCR primer used in the detection method of over-
representation of DNA from male grandparents of the invention
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                                                                                                                                                           Multiplex group PCR primer #239
BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAY-2002; 2002WO-GB002273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-MAY-2001; 2001GB-00011886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ANIM-) ANIMAL HEALTH TRUST
  ABT21492 standard; DNA; 20
                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Swinburne JE
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                                                                                                                                                                                                                                                                                                                                                     WO200292851-A2
                                                                                                                                                                                                                                                                                                  Unidentified
                                                                                                        16-APR-2003
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                                                     ABT21492,
SO CCC CCC CCC X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X
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                                      Gaps
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    0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03; tive 0; Mismatches 1; Indels
Query Match
Best Local Similarity 94.1
Watches 16; Conservative
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594 CCACTGCAAGGTGTACA 610
                       20 ccacrecaacereraca 4
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Novel human protein identification related primer #3.
           20
                                  10-JUL-2003 (first entry)
           ACA90195 standard; DNA;
                       ACA90195
RESULT 1223
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BP.

Human; cytostatic; DAPK3-Agonist; DAPK3-Antagonist; cancer; NOV; PCR;

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primer;
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Bacillus subtilis subsp. subtilis str. 168.

WO2003054163-A2

03-JUL-2003

Homo sapiens

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WO2003031571-A2
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                                                                                                                                                                                                                                                                                            The invention describes an isolated polypeptide comprising any of 33 90-1273 amino acid sequences (1) given in the specification or its mature form, a sequence that is at least 95 % identical to (1), or a sequence comprising one or more conservative substitutions in the amino acid sequence of (1). The polypeptide is useful for preparing a composition for treating or preventing e.g. cancer. This sequence represents a primer used to isolate DNA encoding a novel human NOV protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hyaluronic acid, glycosaminoglycan, hyaluronan synthase, antirheumatic, UDP-glucose 6-dehydrogenase; UDP-glucose pyrophosphorylase; orthopaedic; UDP-N-acetylglucosamine, ophthalmological, dermatcological; joint surgery; eye; rheumatcology; dermatcology; adhesion; development; cell motility; cancer; angiogenesis; wound healing; ss; PCR; primer.
                                                                                                                                                                             New DAPK3 polypeptide, useful for preparing a composition for treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                Kekuda R;
DK, Spytek KA;
         Chaudhuri A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.4; DB 1; Length 20;
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer 11 used to sequence Bacillus subtilis TuaD DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
Alsobrook JP, Burgess CE, Catterton B, June J, Edinger SR, Gerlach VL, Giot L, Gorman L, Guo X, Mezes PS, Millet I, Ooi CE, Patturajan M, Rieger Taupier RJ, Zerhusen BD, Zhong H, Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                  Example 20C; Page 178; 253pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2750
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%;
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ses 16; Conservative
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                                                                                                                                                                                                     preventing e.g., cancer.
                                                                                                                                WPI; 2003-381704/36.
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Matches
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vivlemore401-10.rng

Producing a hyaluronic acid (e.g. for use in eye and joint surgery, orthopedics, rheumatology or dermatology) comprises cultivating a Bacillus host cell and recovering the hyaluronic acid from the

Example 1; Page 37; 218pp; English.

cultivation medium.

Sternberg D,

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Brown

Behr R, Widner W, Tang M, Sternberg D,

WPI; 2003-559139/52.

Sloma A,

(NOVO) NOVOZYMES BIOTECH INC.

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Hyaluronic acid; glycosaminoglycan; hyaluronan synthase; antirheumatic; UDP-glucose 6-dehydrogenase; UDP-glucose pyrophosphorylase; orthopaedic; UDP-N-acetylglucosamine; ophthalmological; dermatological; joint surgery; eye; rheumatology; dermatology; adhesion; development; cell motility; cancer; angiogenesis; wound healing; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer 14 used to sequence Bacillus subtilis TuaD DNA
                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacillus subtilis subsp. subtilis str. 168.
                                                                    Tang M,
                                                                                                                                                                                                                                                                                                                                                                                                                      1288 GTAGCCGTGAAGATGCT 1304
                                               (NOVO ) NOVOZYMES BIOTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABT44021 standard; DNA; 20 BP.
                            21-DEC-2001; 2001US-0342644P.
         20-DEC-2002; 2002WO-US041067.
                                                                   Sloma A, Behr R, Widner W,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-DEC-2001; 2001US-0342644P
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                                                                                                                                                                                                                                                                                                                                                                                                   16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                      WPI; 2003-559139/52.
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Gaps

; 0

1; Indels

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The invention relates to a novel method which comprises producing a hyaluronic acid via cultivating a Bacillus host cell under conditions suitable for production of the hyaluronic acid and subsequently recovering the production of the hyaluronic acid and subsequently recovering the hyaluronic acid from the cultivation medium. The most abundant heteropolysaccharides of the body are the glycosaminoglycans, of which hyaluronic acid is an example. A number of enzymes are involved in the biosynthesis of hyaluronic acid including hyaluronan synthase, UDP-glucose f-dehydrogenase, UDP-glucose pyrophosphorylase and UDP-northy acetylglucosamine. The molecules of the invention demonstrate ophthalmalogical, antirhenmatic and dermatological activities, whilst the method itself may be useful for producing a hyaluronan in a recombinant host cell. The hyaluronan generated may be used in eye and joint surgery, corthopaedics, rheumatology or dermatology and may exhibit further uses within the fields of adhesion, development, cell motility, cancer, angiogenesis and wound healing. The current sequence is that of the PCR primer of the invention which was used during analysis of the enzymes
Producing a hyaluronic acid (e.g. for use in eye and joint surgery, orthopedics, rheumatclogy or dermatclogy) comprises cultivating a Bacillus host cell and recovering the hyaluronic acid from the cultivation medium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      that play a role in the synthesis of hyaluronic acid
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                                                                                                                                                                                                                                                                                                                               Example 1; Page 37; 218pp; English.
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The invention relates to a novel method which comprises producing a hyaluronic acid via cultivating a Bacillus host cell under conditions suitable for production of the hyaluronic acid and subsequently recovering the production of the hyaluronic acid from the cultivation medium. The most abundant heteropolysaccharides of the body are the glycosaminoglycans, of which hyaluronic acid is an example. A number of enzymes are involved in the biosynthesis of hyaluronic acid including hyaluronan synthase, UDP-glucose 6-dehydrogense, UDP-glucose pyrophosphorylase and UDP-cophtalmological, antirheumatic and dermatological activities, whilst the method itself may be useful for producing a hyaluronan in a recombinant host cell. The hyaluronan generated may be used in eye and joint surgery, or the producing and may exhibit further uses within the fields of adhesion, development, cell motility, cancer.

primer of the invention which was used during analysis of the enzymes that play a role in the synthesis of hyaluronic acid

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                                                                                                                                                                                                                                                                                                                                                                                                                        apoptosis; Jagged 2 inhibitor; cytostatic; hyperproliferative disorder;
human; ss; antisense oligonucleotide; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                         Human Jagged 2 chimeric phosphorothioate oligonucleotide SEQ ID NO:51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "phosphorothioate linkages, where all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                    Gaps
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0
                                                                                                                                                                                                                                                                 0.4%; Score 15.4; DB 1; Length 20;
4.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                    Sequence 20 BP; 4 A; 3 C; 5 G; 8 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "2'-O-methoxyethyls"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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'mod base= OTHER
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                                                                                                                                                                                                                                                                                                  1288 GIAGCCGTGAAGATGCT 1304
                                                                                                                                                                                                                                                                                                                                                          ADD00996 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                          94.1%;
                                                                                                                                                                                                                                                                                                                  4 GTAGCCGTTAAGATGCT
                                                                                                                                                                                                                                                                                                                                                                                        01-JAN-2004 (first entry)
                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 94.1'
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                         2'-O-methoxyethyl.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                          ADD00996;
                                                                                                                                                                                                                                                                                                                                         RESULT 1226
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The present invention describes a method of determining whether an individual is predisposed to susceptibility to low bone mineral density (BMD) and/or bone damage comprising identifying whether the individual has at least one polymorphism in a polymucleotide encoding a protein, where the polymucleotide is one of 81 200-500 nucleotide sequences (S1, see ADC98235 to ADC98135). An agent identified in an method from the present invention which can be used for the prevention or treatment of a disease resulting in susceptibility to low BMD and/or bone damage is usceptibility to low BMD and/or bone damage is susceptibility to low BMD and/or bone damage is susceptibility to low BMD and/or bone damage. The disease associated with low BMD and/or bone damage is osteoporosis. The present PCR primer sequence is used in the exemplification of the present invention.
                                                                                                                                                                    Determining whether an individual is predisposed to susceptibility to low bone mineral density (BMD) and/or bone damage, involves identifying polymorphisms in associated genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a tubercle bacillus gene chip and its application. According to the characteristics of tubercle bacillus genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                             Bennett S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A tubercle bacillus gene chip useful for tuberculosis diagnosis and reasonable selection of medicine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.4; DB 1; Length 20; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                             Mangion J, Galwey N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 13 A; 1 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             subercle bacillus gene chip; ss; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 61; 4pp; Chinese.
                                                                                                                                                                                                                                                               Example 8; Page 239; 246pp; English
                                                               Townley DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Probe related to the invention #61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JAN-2001; 2001CN-00107010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%;
                    (INCY-) INCYTE GENOMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3117 TTAATTTTTAACTTATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18 TTAATTTTAATTTATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADF73003 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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Best Local Similarity 94.1
Matches 16, Conservative
                                                             Valdes A,
Schafer A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-240333/24.
                                                                                                                               WPI; 2003-559156/52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BAOL/) BAO L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                               Jones KA,
                                                                                      Mckay I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF73003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bao L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1228
셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method for inducing apoptosis in a cell inhibitor to reduce Jagged 2 levels or activity. Also described: 1 treating a subject having a disease or condition associated with insufficient apoptosis by administration of a Jagged 2 inhibitor; (2) a pharmaceutical composition comprising a Jagged 2 inhibitor and another active ingredient for inducing apoptosis; and (3) a kit comprising a the Jagged 2 inhibitor in the Induction of apoptosis; The Jagged 2 inhibitor in active ingredient for inducing apoptosis; and (3) a kit comprising a the Induction of apoptosis. The Jagged 2 inhibitor in activity. The method can be used for inducing apoptosis in a cell or animal for treating a subject having a disease or condition associated animal for inducing a subject having a disease or condition associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               low bone mineral density; BMD; bone damage; polymorphism; osteoporosis; single nucleotide polymorphism; SNP; PCR primer; ss; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence represents a human Jagged 2 chimeric phosphorothioate antisense oligonucleotide, which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                               a subject having administering to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                with insufficient apoptosis, e.g., hyperproliferative disorder. The
                                                                                                                                                                                                                                                                                                                          Inducing apoptosis in a cell or animal for treating a subject har condition associated with insufficient apoptosis by administering cell or animal a Jagged 2 inhibitor to reduce Jagged 2 levels or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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Pred. No. 1.4e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OMD_01 polymorphism marker PCR primer S primer seq.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 15; SEQ ID NO 51; 148pp; English
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                                                                                                                                                    12-MAR-2002; 2002US-00096399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%;
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04-NOV-2002; 2002US-0423559P.
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                                                                                                        10-MAR-2003; 2003WO-US007340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CACCTGCAAGGACCTGG
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les 16; Conservative
                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                          Shapard PJ;
                                                                                                                                                                                                                                                                                      WPI; 2003-756943/71
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                      WO2003077848-A2
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                                                                  25-SEP-2003
                                                                                                                                                                                                                                             Koller E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
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Best Local S:
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                               activity.
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RESULT 1227 ADC98542,

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sequence and molecular mechanism produced by tubercle bacillus resistance to drug, and according to its application the probe can be designed and selected, and the probe array can be regularly and reasonable arranged according to a certain mode to form optimized probe array. The gene chip can be used for quickly, accurately and high-efficiency identifying tubercle bacillus, and can be used for screening and detecting its resistance to drug, and can detect the mutation of specific site of tubercle bacillus gene sequence and the mutation of non-specific site. The invention can be used for tuberculosis diagnosis and reasonable selection of medicine. The present sequence represents a probe related to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Phosphorothioate backbone where all cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides for modulating Jagged 2 expression, useful for diagnosing, preventing or treating diseases or conditions associated with Jagged 2, e.g. cancer or developmental disorders.
                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense; Jagged 2; hyperproliferative disorder; cancer; developmental disorder; apoptosis; prophylaxis; antisense therapy; phosphorothioate; human; ss.
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/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                     Query Match 0.4%; Score 15.4; DB 1; Length 20; Best Local Similarity 94.1%; Pred. No. 1.4e+03; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Jagged 2 antisense oligonucleotide ISIS #148740.
                                                                                                                                                                                         Sequence 20 BP; 3 A; 9 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
                                                                                                                                                                                                                                                                                   836 TGGTGGTGCTGCCAGCC 852
                                                                                                                                                                                                                                                                                                                                                                                           ADH62956 standard; DNA; 20 BP
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                                                                                                                                                             the invention.
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                          The invention relates to novel antisense compounds targetted to a nucleic acid molecule encoding Jagged 2 to inhibit its expression. Antisense compounds of the invention are useful for treating an animal having a disease or condition associated with Jagged 2, e.g. hyperproliferative disorder (particularly cancer), a developmental disorder or a disease or condition that arises from aberrant apoptosis. They are also used for diagnostics, prophylaxis or as research reagents or kits. The invention is also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human Jagged 2 DNA. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides of 8-40 nucleobases, useful for modulating
the function of nucleic acid molecules encoding Jagged 2, ultimately
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Phosphorothioate antisense DNA oligo to modulate human Jagged 2 SeqID 51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; ss; antisense; Jagged 2; differentiation; cell fate; signalling;
Usher syndrome type 1a; retinitis pigmentosa; phosphorothioate backone;
2' MOE wing.
                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                 0.4%; Score 15.4; DB 1; Length 20;
44.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1. .20
//tag= b
/mote= "OTHER phosphorothioate backbone"
                                                                                                                                                                                                     Seguence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;
Claim 3; SEQ ID NO 51; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                  1542 CACCTTCAAGGACCTGG 1558
                                                                                                                                                                                                                                                                                                                                                                                                              BP
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                                                                                                                                                                                                                                                   94.18;
                                                                                                                                                                                                                                                                                                                    17 CACCTGCAAGGACCTGG
                                                                                                                                                                                                                                                                                                                                                                                                            ADH57111 standard; DNA; 20
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/*tag= a
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                                                                                                                                                                                                                                                 Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2003207839-A1.
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modified_base
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Homo sapiens.
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This invention relates to movel antisense compounds that can be used to modulate the expression of Jagged 2. Specifically, it refers to compositions useful for inhibiting the expression of Jagged 2, a human homologue of the Drosophila Serrate gene, which is involved in differentiation and cell fate, as well as positive feedback control over signalling genes such as Notch 1, Notch 3 and Jagged 1. The Jagged 2 gene is located on chromosome 14932, a region that has been implicated in retinitis pigmentosa. The present invention describes antisense oligonucleotides that comprise at least one modified sugar moiety, a 2'-O methoxyethyl (2' MOE) and at least one modified sugar moiety, a 2'-O methylcytosine. These compounds are useful for modulating the function of nucleic acid molecules encoding Jagged 2, ultimately modulating the reagents and in diagnostics. This oligonucleotide sequence is a smount of Jagged 2 produced, which in turn is useful for research reagents and in diagnostics. This oligonucleotide sequence is a phosphorothioate antisense DNA oligo used to modulate human Jagged 2 expression in an exemplification of the invention.
modulating the amount of Jagged 2 produced.
                                                      Claim 3; SEQ ID NO 51; 63pp; English
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Sequence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;

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0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03;
                              1; Indels
                              0; Mismatches
                                                           1542 CACCTTCAAGGACCTGG 1558
                                                                           17 CACCTGCAAGGACCTGG
                              16; Conservative
                Local Similarity
    Query Match
                                 tches
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Human PDE4C oligonucleotide sequence. ABZ99247 standard; DNA; 20 BP (first entry) 17-OCT-2003 ABZ99247; RESULT 1231
ABZ99247/C
XX
AC ABZ99247/C
DT 17-OCT-5
XX
M Human; FW
M antiastl
XW
Antiastl
Antiastl
XW
Antiastl
Antias

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiathmatic; hypotensive; immunosuppressive; cycostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P.

Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, I Miller S,

(EPIG-) EPIGENESIS PHARM INC

ä

Aguilar

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 14489; 872pp; English.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypoteensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antialmanatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing benefit in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fig. at ftp.wipo.int/pub/published_pot_sequences
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Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;

Gaps ö Score 15.4; DB 1; Length 20; Pred. No. 1.4e+03; Indels 0; Mismatches 0.4%; Best Local Similarity 94.1 Matches 16; Conservative Query Match

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1884 CTTCAAGCTGCTGAAGG 1900 20 cricaAccieciecaes

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Gaps

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ABZ91972 standard; DNA; 20 17-OCT-2003 (first entry) ABZ91972; ABZ91972,

B

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiinsthmatic; hypotensive; fumunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

sapiens. Ношо WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar Katz E, Pabalan J, Nyce JW, Li Y, Sandrasagra A, Ke Miller S, Tang L, Shahabuddin S;

ä

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 7214; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding regaion, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions. of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entilnflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entilating manacory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfaction in a subject, arearisating bronchodilation, arearisating hypotensity, lung in a subject of an entiled and a subject of an entiled and a subject of a lung surfaction in a subject of a lung as unfaction in a subject of a lung surfaction in a subject of a lung as unfaction in a subject of a lung as unfaction in a subject of a lung surfaction in a subject of a lung as unfaction in lung as lung and a lung and lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 12 A; 3 C; 0 G; 3 T; 0 U; 2 Other;

0.4%; Score 15.4; DB 1; Length 20; 84.2%; Pred. No. 1.4e+03; 3; Indels 0; Mismatches 3316 TTTAGGAGATTTATTTT 3334 20 TTTAGGAGANTTTTNTTT 2 16; Conservative Local Similarity Query Match Matches 엄 à

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Gaps

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ABD32278 standard; DNA; 20 BP

29-JUL-2004 (first entry)

ABD32278;

Human PDE4C-derived oligonucleotide SEQ ID 14489.

beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; ABD3278/
XX ABD3
XX ABD3
XX ABD3
XX ABD3
XX Huma
XX Huma
XX Huma
XX Huma
XX Huma
XX ABD1
XX ABD1
XX ABD1
XX WO2C
XX BD1
XX ABD1
XX ABD

Homo sapiens

WO200285309-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Katz E, Pabalan J, s; Sandrasagra A, L, Shahabuddin Li Y, San , Tang L, Miller S, Ę, Nyce

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

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This invention describes a nowest composition, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating cypression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery cypression of the invention has antiallergic, antiinflammatory, antiasthmatic, instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, whypertension, emphysema, chronic observations, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the targ
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                                                                                                 invention describes a novel composition (a) a first active agent,
                            Claim 15; SEQ ID NO 14489; 763pp; English.
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Gaps . 0 0.4%; Score 15.4; DB 1; Length 20; 44.1%; Pred. No. 1.4e+03; ve 0; Mismatches 1; Indels Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other; 94.18; Matches 16; Conservative Local Similarity Query Match

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1884 CTTCAAGCTGCTGAAGG 1900

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20 CTTCAAGCTGCTGCAGG

ABD28202 standard; DNA; 20 BP. ABD28202; ABD28202,

(first entry) 29-JUL-2004

R19956-derived oligonucleotide SEQ ID 7214.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antialergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; abeta-adrensici agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

/mod_base= OTHER /note= "OTHER= Phosphorothioate backbone. All cytidines

are 5-methylcytidines"

1. .20 /*tag=

l. .5 /*tag= a

/mod_base= OTHER /note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"

/mod_base= OTHER /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"

/*tag= c

15. .20

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New antisense oligonucleotides for modulating LIM domain kinase 1 expression, useful for diagnosing, preventing or treating conditions associated with the kinase, e.g. neurological or developmental disorders.
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                                                                                                                                                                                                                                                                                                                                                                          18-JUL-2002; 2002US-00199199
                                                                                                                                                                                                                                                                                                                                       .8-JUL-2002; 2002US-00199199
                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VPI; 2004-121553/12.
                                                                                                                                                                                                                                                               US2004014047-A1
modified base
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                                                                                         modified base
                                                                                                                                                                                                                                                                                                    22-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cowsert LM,
This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating construction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition is useful for preventing or comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung cinflammation, allergies and/or bronchoconstriction and/or lung distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary chartension, emphysema, chronic obstructive pulmonary disease, pulmonary intertions brain or election, pulmonary intections, brained adenosine content of the anti-sense object the pulmonary the breakdown of the reduced adenosine content of the anti-sense to prevent the breakdown of the oligonucleotides into products that free adenosine into the preduced adenosine content of the anti-sense operonment and thereby to e.g. lung. brain, heart, kidney, etc. tissue environment and thereby to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                      Katz E, Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 12 A; 3 C; 0 G; 3 T; 0 U; 2 Other;
                                                                                                                                                                                                                                     Claim 15; SEQ ID NO 7214; 763pp; English.
                                                    Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                 (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                      bronchodilating agent.
                                                                                                             WPI; 2003-093058/08
                                                    Nyce JW, I
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The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding LIM domain kinase 1. The compound operationally hybridises with the nucleic acid molecule encoding LIM domain kinase 1 and inhibits the expression of LIM domain kinase 1. It specifically hybridises with at least an 8-nucleobase portion of a preferred trager region on the nucleic acid molecule encoding LIM domain kinase 1. The antisense oligonucleotied is useful for modulating the expression of LIM domain kinase 1 in cells or tissues to treat diseases associated with their expression, such as a developmental disorder or a neurological disorder. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neuroprotective, LIM domain kinase 1, developmental disorder, neurological disorder, diagnostic, prophylaxis, human, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.4; DB 1; Length 20; 34.1%; Pred. No. 1.4e+03; ve 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                        human LIM domain kinase 1 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
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nes 16; Conservative
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.4; DB 1; Length 20;
14.2%; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human LIM domain kinase 1 antisense oligonucleotide #83.
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20 TTTAGGAGANTTTTNTTTT 2

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84.2%;

16; Conservative

Matches

Best Local Similarity

Query Match

ADI38799 standard; DNA; 20 BP

(first entry)

22-APR-2004

RESULT 1235
AD138799/C
XX
XX
AC AD138799
AC AD138799
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DT 22-APR-2
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M neuropro
XX
M neurolog
XX
COS Homo sag
XX
FH Key

ADI38799;

Homo sapiens

Location/Qualifiers

Homo sapiens

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Gaps ö

vivlemore401-10.rng

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotides for modulating LIM domain kinase 1 expression, useful for diagnosing, preventing or treating conditions associated with the kinase, e.g. neurological or developmental disorders.
                                                /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                 /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                     /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human LIM domain kinase 1 antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 15; SEQ ID NO 21; 81pp; English.
                                                                                  are 5-methylcytidines"
Location/Qualifiers
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/*tag= a
/mod_base= OTHER
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/mod_base= OTHER
.^mueR= 2'
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                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                        Cowsert LM, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-121553/12.
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                                                                                                                                                                                                                                                      US2004014047-A1.
Key
modified_base
                                                                                                 modified_base
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ADJ61132/c
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WWW.XEXPXPXPX IN
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Novel single or multiple target oligonucleotide anti-sense to e.g. inititation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.

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Aguilar D, Miller

Nyce JW, Tang L, Sandrasagra A, Shahabuddin S, Lu H, Cong H;

WPI; 2004-203534/19.

29-JUL-2002; 2002US-0399076P. 25-JUL-2003; 2003WO-US023509.

WO2004011613-A2,

05-FEB-2004

Homo sapiens.

(EPIG-) EPIGENESIS PHARM INC.

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The present invention relates to an oligonuclectide anti-sense to e.g., initiation codon, coding region with 2-10 nuclectides of 5'-end and 3'-end of nucleic acid target comparising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the cligonuclectide and optionally surfactant operatively linked to the oligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(iss), asthma, impeded chosen from airway inflammation, allergy(iss), asthma, impeded conspiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway constitution. The present sequence represents an oligonuclectide of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; SEQ ID NO 1988; 85pp; English.
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modified base
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vivlemore401-10.rng

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New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding resistin, useful for treating a metabolic disorder,
e.g. diabetes or obesity, or atherosclerosis.
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                        Phosphorothioate backbone. All cytidines
                                                                                                                                                                                                          /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                            /*tag= a
/mod_base= OTHER
/note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
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Pred. No. 1.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 5 A; 7 C; 2 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 50; 75pp; English.
                        /note= "OTHER= Phospho
are 5-methylcytidines"
  /mod_base= OTHER
                                                                                                                                                                                    /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1352 TGGAGATGATGAAGATG 1368
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Best Local Similarity 94.1%;
Matches 16; Conservative
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                                                                                                                                                              15. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Freier SM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK79424;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding resistin, useful for treating a metabolic disorder,
e.g. diabetes or obesity, or atherosclerosis.
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                     /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                                     /*tag= a
/mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
15. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   represents a human resistin antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human resistin antisense oligonucleotide seq id 50.
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                                                                                                                                                                                                                                     base= OTHER
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ADJ38661 standard; DNA; 20 BP
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modified base
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AC AbJ3
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Gaps

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                                                                                                                                                   New antisense compound targeted to a nucleic acid molecule encoding about.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 11 A; 2 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 6758; 417pp; English
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                14-AUG-2002; 2002US-0403416P.
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                                              (PHAA ) PHARMACIA CORP
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                                                                                                                    WPI; 2004-203785/19
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                                                                                    Roberds SL;
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ADK79423/c
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New antisense compound targeted to a nucleic acid molecule encoding

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                                                                                                                          nucleic acid molecule encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligomucleotide with 2. MOE wings and a deoxy gap. Used during the antisense inhibition of human Navi.3 expression, the oligomucleotides are designed to target different regions of the human Navi.3 RNA.
Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                 invention relates to an antisense compound targeted to
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4.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 13 A; 2 C; 2 G; 3 T; 0 U; 0 Other;
                                                                            Claim 4; SEQ ID NO 6757; 417pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    ilarity 94.1%;
Conservative
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nes 16; Conserv
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12-OCT-2001; 2
15-OCT-2001; 2
17-OCT-2001; 2
22-OCT-2001; 2
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                                           Query Match
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(GERL/)
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associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeninal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder; including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to an antisense compound targeted to a mucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate bain from burns, migraine headache, cluster headache, mild-to-moderate but not limited to neonatal or infantile epilepsy, or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide, with luman Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                       Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                              Chimeric phosphorothioate oligonucleotide to target Navl.3 #7215.
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                                                                                                                                                                       0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03;
                                                                                                                                                                                                1; Indels
                                                                                                                                               Sequence 20 BP; 12 A; 2 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 7215; 417pp; English.
                                                                                                                                                                                                                           3260 GATATTTTATTTGCTTT 3276
                                                                                                                                                                                                                                                                                                                   ADK79881 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-AUG-2002; 2002US-0403416P.
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                                                                                                                                                                                   94.1%;
                                                                                                                                                                                                                                                  19 GATATTTTATTTGCCTT
                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                      20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Roberds SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                           ADK79881;
                                                                                                                                                                         Query Match
                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                         RESULT 12.
ADK79881/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic; antidiabetic; anorectic; cerebroprotective; neuroprotective; antinflammatory; thyromimetic; gene therapy; antisense therapy; NOVX polypeptide related disorder; cancer; diabetes; obesity; endocrine disorder; CNS disorder; inflammatory disorder; chromosome mapping; tissue typing; predictive medicine; intracellular protiein-like protein; sorting nexin 6-like protein; 573045109RK cyclin-like protein; cMob5 cancer specific protein; phosphatidyjethanolamine-binidng protein-like protein;
                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoglobulin-like LRR-domain containing protein;
NUMB binding protein LNXp80-like protein;
zinc finger protein-like protein;
actin-binding protein alpha-like protein;
actin-binding protein frabin-alpha-like protein;
actin-bending protein frabin-alpha-like protein;
hapatocallular carcinoma autoantigen-like protein;
haematopoietic stem/progenitor cells protein MDS029-like protein;
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                                                     0.4%; Score 15.4; DB 1; Length 20;
14.1%; Pred. No. 1.4e+03;
ve 0; Mismatches 1; Indels
Sequence 20 BP; 12 A; 2 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human protein Nov3 probe segid 72
                                                                                                                                                                                          3260 GATATTTTATTTGCTTT 3276
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2001US-0328449P
2001US-0339414P
2001US-0330142P
2001US-0341658P
2001US-0343629P
2001US-03436357P
                                                                                                                                                                                                                                                                                                                                                                                                                  ADO09378 standard; DNA; 20 BP.
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2001US-0327917P.
2001US-0328029P.
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                                                                                         94.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-2004 (first entry)
                                                                                         Local Similarity 94.1
les 16; Conservative
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vivlemore401-10.rng

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; imPGES-1; mPGES-1 inhibitor; inhibitor; categlandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; catediant; neuroprotective; antiinflammatory; neuroprotective; austrintic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-SEP-2003; 2003WO-US030374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-SEP-2002; 2002US-0413549P.
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                                                                                                                                                                                                                                                                 Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                   modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gierse JK;
                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ischemia.
                                                                                                                                                                                                                    Homo
 defined sequence of, a mature form, one or more conservative cubstitutions or at least 95% identity to 158 amino acids SEQ ID NO: 2n substitutions or at least 95% identity to 158 amino acids SEQ ID NO: 2n cas given in the specification, where n is an integer between 1-33. The novel proteins of the invention are members of the following families: novel proteins of the invention are members of the following families: intracellular protein-like proteins; sorting nexin 6-like proteins; cMob5 cancer specific proteins; ERP16 protein-like proteins; cMob5 cancer specific proteins; ERP16 proteins; cMob5 cancer specific proteins; ERP16 proteins; cMob5 cancer specific proteins; ERP16 proteins; containing proteins; actin-binding proteins; actin-binding protein alpha-like proteins; actin-binding protein frabin-alpha-like proteins; actin-binding protein specific proteins; actin-binding proteins; haematopojetic contoining in the proteins; and tumour protein MDS029-like proteins; TRAP-delta-like proteins; and tumour protein MDS029-like protein; TRAP-delta-like proteins; and tumour protein p53-binding protein 130-like proteins; carrivin light chain-like proteins; leucine-rich protein; and tumour protein p53-binding protein 2-like proteins; carrivin of the Novx polypeptide, such as cancer, diabetes, obesity, and endocrine, CNS and inflammatory disorders. They can also be used in various detection and screening assays, chromosome mapping, tissue typing and predictive medicine. This sequence represents a probe used in the carrivin of DNA encoding a novel human polypeptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                  New isolated NOVX polypeptides and nucleic acid molecules useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, such as cancer, obesity, diabetes and inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention describes a new isolated polypeptide (I) comprises a fully
                                                                                                                                                                                                                                                 Catterton E, Chant JS, Chaudhuri A;
: L, Gorman L, Guo X, Kekuda R, Mezes P?
jan M, Rieger DK, Spytek KA, Taupier RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 20C; SEQ ID NO 72; 164pp; English.
                                                                                                                                                                                                                                                Alsobrook JP, Burgess CE, Cattert
Edinger S, Gerlach V, Giot L, Go
Millet I, Ooi CE, Patturajan M,
                                                                                                                                                                                                                                                                                                   Zhong M;
                                                                                                                                                                                                                                                                                                  Zhong H,
                                                              MEZES P S.
MILLET I.
OOI C E. BATTURAJAN M.
RIEGER D K.
SPYTER K A.
TAUPIER R J.
ZERHUSEN B D.
                                                                                                                                                                                                                                                                                                                                 WPI; 2004-108207/11
GIOT L.
GORMAN L.
                                                                                                                                                                                                 ZHONG H. ZHONG M.
                                                KEKUDA R
                                                                                                                                                                                                                                                                                                                                                                                                                    CNS diseases.
                                                                                                                                                                                                                                                                                                   Zerhusen BD,
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                                                                                                                                                                                (ZERH/)
(ZHON/)
(ZHON/)
                                                                               (MILL/)
(OOIC/)
(PATT/)
(RIEG/)
                                                                                                                                                 (SPYT/)
(TAUP/)
                                                                 MEZE/)
                                                KEKU/)
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/mod_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

Location/Qualifiers

sapiens.

1. .20 /*tag=

/*tag= a /mod_base= OTHER /note= "2'-0-methocyethyls" 16. .20

note= "2'-0-methoxyethyls"

/*tag= c /mod_base= OTHER

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mDGES-1). The human mpcRES-1 gene is located on chromosome 9, more specifically to olystopic mpcRES-1 gene is located on chromosome 9, more specifically to olystopic mpcRES-1 and sequence comprising 8-30 bp targeted to a nucleic acid encoding mpcRES-1, which specifically whybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpcRES-1 in cells or tissues; and (3) a method of treating an animal condition associated with mpcRES-1. MPCRES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamatory, neuroprotective, nootropic, antidiamatory, neuroprotective, nootropic, antidiamatory, usaotropic and cardiovascular activities, and can be used as mpcRES-1 inhibitors and in gene therapy. The antisense compound to be used as mpcRES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpcRES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or copithalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 8 A; 8 C; 2 G; 2 T; 0 U; 0 Other;
Claim 4; SEQ ID NO 1171; 132pp; English
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2750 ო

2734 TACCTGAAGATGGGAGC 19 raccregagaregage

8 셤

Local Similarity 94.1 nes 16; Conservative

Best Loca Matches

ADM14984 standard; DNA; 20

RESULT 1245

01-JUL-2004 (first entry)

ADM14984;

ADM14984/c ID ADM149 XX AC ADM149 XX DT 01-JUL

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sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                                                          chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; mucrosomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGBS-1). The human mPGBS-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                     Gaps
                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1204.
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0
  Length 20;
                    1; Indels
0.4%; Score 15.4; DB 1;
94.1%; Pred. No. 1.4e+03;
cive 0; Mismatches 1;
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/note= "2'-O-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                         2332
                                                                                                               BP.
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                                         2316 TCTGTGTGTGTGTGTGT
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                                                                                                                                                       (first entry)
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 Query Match 0.4
Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-305094/28.
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                                                                                                                                                                                                                                                                                                                                   Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                              modified base
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                                                                                                                                                                                                                                                                                                      sapiens
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                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gierse JK;
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                                                                                                                                   ADM15017;
                                                                                           RESULT 1246
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                                                                                                     ADM15017,
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mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamatory, neuroprotective, antidiamatory, neuroprotective, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disoacer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; imPGES-1; mPGES-1 inhibitor; intidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; assotropic; ophthalmological; immunomodulatory; cardiavascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
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/note "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 8 A; 8 C; 3 G; 1 T; 0 U; 0 Other;
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nes 16; Conservative
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004028458-A2
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modified_base
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ADM15000/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          셤
                                                                                                                                                          The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGBS-1). The human mpGBS-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-10 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibite its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal harisense condition associated with mpGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthitic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemmia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
                                                          New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 1.4e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 20 BP; 9 A; 8 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                 Claim 4; SEQ ID NO 1595; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2334 CGTGTGTGTGTGTGT 2350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM15146 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 cererererererer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Д
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 94.1
Matches 16; Conservative
                             WPI; 2004-305094/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
 Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                     ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM15146;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1248
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8
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mGCBS-1). The human mmcGBS-1 gene is located on chromosome 9, more specifically to ogd4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding meDGBS-1, which specifically hybridise with the nucleic acid encoding compounds inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal carbonial adisease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, assotropic, antidiamaclosy, neuroprotective, nootropic, antidiamaclosy, neuroprotective, nootropic, antidiamaclosy, or sphthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used as wed for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alabeimer's diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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Pred. No. 1.4e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 9 A; 7 C; 2 G; 2 T; 0 U; 0 Other;
'note≈ "2'-0-methocyethyls"
                                                                                           /mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 1333; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2350
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                                                                                                                                                                                                                                                                                                                               25-SEP-2003; 2003WO-US030374.
                                                                                                                                                                                                                                                                                                                                                                                             25-SEP-2002; 2002US-0413549P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2334 CGTGTGTGTGTGTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 cererererererer
                                                                    U
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                           16. 20
                                                                *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-305094/28.
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                                                                                                                                                                                              WO2004028458-A2
                              modified base
                                                                                                                                                                                                                                                               08-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM15000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; Eccaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory diserses syndrome; pulmonary phyertenation; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.

25-JUL-2003; 2003US-00627930. 23-APR-2002; 2002WO-US013135. 23-APR-2002; 2002WO-US013143.

US2004049022-A1.

11-MAR-2004.

Homo sapiens.

SANDRASAGRA A.

NYCE J W.

NYCE/)

SHAHABUDDIN S.

(MILL/) (SHAH/) (LUHH/) CONG/)

CONG H.

AGUILAR D.

TANG L.

TANG/) AGUI/)

MILLER S.

ADO46522 standard; DNA; 20 BP

RESULT 1250 ADO46522/c

Human oligonucleotide #1888.

(first entry)

15-JUL-2004 ADO46522;

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/note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.4; DB 1; Length 20;
94.1%; Pred. No. 1.4e+03;
ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 8 A; 7 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                 note = "2'-0-methoxyethyls"
                                                                                                                                /note= "2'-0-methocyethyls"
16. .20
                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 1187; 132pp; English,
                                                       Location/Qualifiers
                                                                                 base= OTHER
                                                                                                                        /mod_base= OTHER
                                                                                                                                                  /*tag= c
/mod base= OTHER
                                                                                                                                                                                                                 25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                  25-SEP-2002; 2002US-0413549P.
                                                                                                                  ಗ
                                                                         Д
                                                                                                                 /*tag=
                                                                        *tag=
                                                                                                                                                                                                                                                 (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                  VPI; 2004-305094/28
                                                                                                                                                                                  WO2004028458-A2.
                                                        Key
modified_base
                                                                                                         modified base
                                                                                                                                         modified base
                                 Homo sapiens.
Synthetic.
                                                                                                                                                                                                  08-APR-2004.
                                                                                                                                                                                                                                                                   Gierse JK;
                                                                                                                                                                                                                                                                                                                            ischemia.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to oppose 1 and 1 antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisflammatory, neuroprotective, antisflammatory neuroprotective, neuroprotective, antisflammatory ineuroprotective, antisflammatory ineuroprotective, antisflammatory ineuroprotective, antisflammatory and and in gene therapy. The antisense compound be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.; inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion inhury, or ophthalmic, immunological, cardiovascular or neurological disorder.
New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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L, Aguilar D, Miller

Nyce JW, Sandrasagra A, Tang Shahabuddin S, Lu H, Cong H;

WPI; 2004-293804/27.

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The invention relates to oligonuclectides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nuclacidese of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL) chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL) crossen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL) crosses a, tryptase b, pDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention and/or tractment of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the construction and/or tractment of a respiratory or lung disease. The coligonucleotides are useful for reducing or inhibiting expression of a construction and/or tracting a respiratory or lung disease. The creptorides b, PDE4 B, PDE4 C, or PDE3, ICAM, VCAM, tryptase a, creptoratory or lung disease. The creptorides b, PDE4 B, PDE4 C, or PDE3, ICAM, VCAM, tryptase a, creptoratory or lung disease. The creptoride, and/or asthma and/or lung allergies associated with creceptor(s), chronic obstructive pulmonary disease (COPD), creceptor fibrosis (CF), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory distances syndrome, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel single or multiple target oligonuclectide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; SEQ ID NO 1988; 174pp; English.
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Gaps

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2334 CGTGTGTGTGTGTGT 2350

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CGTGTGTGTATGTGT

18

94.18;

16; Conservative

Best Local Similarity Matches 16, Conserv

Query Match

ó

Gaps

.; 0

Length 20; Indels

0.4%; Score 15.4; DB 1; 94.1%; Pred. No. 1.4e+03; live 0; Mismatches 1;

Best Local Similarity 94.1 Matches 16; Conservative

Query Match

3713 CAGAGGTGTCACCCAAA 3729

8 8

2 CAGAGCTGTCACCCAAA 18

Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;

given in table 5 of the invention.

SXS

vivlemore401-10.rng

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This invention relates to the identification of a novel susceptibility locus AST-1 for asthma and other IgE mediated diseases mapped to the human chromosome 7014-p15. Specifically, it refers to two overlapping can ensure an amely GPRA (G-protein coupled receptor for asthma susceptibility) and AAA1 (asthma associated alternatively spliced gane 1). The present invention describes identifying single nucleotide polymorphisms, as well as insertion or deletion polymorphisms, occurring at different positions in the AST-1 locus, and furthermore providing vectors, host cells, primers and probes in order to determine the status of an individual. Accordingly, it provides a kit to diagnose or assess predisposition to cathma, chronic obstructive pulmonary disease or cancer and other IgE mediated disease including rhinitis and dermatitis, such that derived pharmaceutical compositions exhibit cytostatic and antiasthmatic activities. Purthermore, it provides a transgenic animal comprising the asthma locus-1 (AST-1) DNA. This obligonucleotide sequence is a PCR primer used to sequence the exon and exon/ intron boundaries of human GPRA DNA,
                                                                                                                                                                 ö
hypertension, lung inflammation, bronchitis, airway obstruction or
bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New GPRA polypeptides, useful in preparing a composition for diagnosing, treating or preventing asthma, other IgE-mediated disease, chronic obstructive pulmonary disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vendelin J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fwd PCR primer used for sequencing exon 9b boundary of human GPRA DNA
                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; AST-1; asthma; IgE mediated disease; human; GPRA; G-protein coupled receptor for asthma susceptibility; AAA1; asthma associated alternatively spliced gene 1; primer; PCR; chronic obstructive pulmonary disease; cancer; rhinitis; dermatitis; cytostatic; antiasthmatic; transgenic; asthma locus-1.
                                                                                                                                                               ;
                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polvi A, Maekelae S,
                                                                                                                                                             1; Indels
                                                                             Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                     Score 15.4; DB 1;
Pred. No. 1.4e+03;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Laitinen T, Kere J, Laitinen LA,
Pulkkinen V, Salmikangas P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 7; Page 76; 265pp; English.
                                                                                                                                                                                                     1884 CTTCAAGCTGCTGAAGG 1900
                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-DEC-2002; 2002US-0435846P.
03-JAN-2003; 2003US-0437895P.
26-MAR-2003; 2003US-048767P.
09-UUL-2003; 2003US-0486000P.
                                                                                                                                                                                                                                          20 crrcaagcrecrecage 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-DEC-2003; 2003WO-FI000973.
                                                                                                                     Query Match 0.4%;
Best Local Similarity 94.1%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                             ADP84328 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-500286/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                             23-SEP-2004
                                     invention.
                                                                                                                                                                                                                                                                                                                                                                                       ADP84328;
                                                                                                                                                                                                                                                                                                         RESULT 1251
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883333
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         extracellular response kinase (MAPK/ERK) kinase kinase (MEKK), specifically designated WEKK1, MEKK2 and MEKK3. The MEKK proteins are used to modulate and regulate signal transduction in cells, as well as for regulation of gene transcription in a cell encoding MEKK, where the cell is involved in inflammation, regulation of cellular proliferation and differentiation, regulation of development, regulation of cell death or regulation of inflammation. They are also used to prepare antibodies. MEKK polynucleotides can be used to produce the protein recombinantly and as a source of probes and primers. The present sequence represents a PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes human mitogen-activated protein kinase/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human MEKK polynucleotides and polypeptides, used for regulating signal transduction in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for human MEKK2, which is used in an example from the present
                                                                                                                                                                 MEKK1; MEKK2; MEKK3; mitogen-activated protein kinase; MAPK; ERK extracellular regulated kinase; signal transduction; regulation; MARK;ERK; MEK; MKKK; inflammation; cellular proliferation; differentiation; development; cell death; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 15.4; DB 1; Length 21; 94.1%; Pred. No. 1.5e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 5 A; 3 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                   Human MEKK2 PCR primer SEQ ID NO:28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 64; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001 GCAGCTGGTGGAGGACC 2017
                                 AAZ25089 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                        98US-0078153P.
98US-0099165P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 ggagcregregagec 17
                                                                                                                                                                                                                                                                                                                                                                        99WO-US005556.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CADU-) CADUS PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-571843/48.
                                                                                                                                                                                                                                                     Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                       WO9947686-A2
                                                                                                                                                                                                                                                                                                                                                                        15-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                        16-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                         04-SEP-1998;
                                                                                                  09-DEC-1999
                                                                                                                                                                                                                                                                                                                                       23-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Johnson GL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer
RESULT 1252
AAZ25089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loc
Matches
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New polynucleotide encoding a phosphatidic acid-preferring phospholipase Al polypeptide, useful for treating e.g. CNS, genitourinary, cardiovascular or hematological disorders, and asthma or cancer.
                                                                                                                                                 haematological disorder; phosphatidic acid-preferring phospholipase A1; CNS; genitourinary; asthma; chronic obstructive pulmonary disease; cancer; enzyme; gene therapy; human; PCR; primer; ss.
                                                                                                                                   Cytostatic; antiasthmatic; neuroprotective; cardiant; cardiovascular;
                                                                                      Phosphatidic acid-preferring phospholipase Al PCR primer 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 61; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                   18-FEB-2002; 2002WO-EP001684.
                                                                                                                                                                                                                                                                                                                                                                                                                             21-FEB-2001; 2001US-0269856P.
29-JUN-2001; 2001US-0301483P.
                                       19-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-674934/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                            WO200266623-A2.
                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                      29-AUG-2002.
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  AAL41510;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method for inducing an immune response against B-lymphoproliferative disorders. The method comprises DNA sequences encoding for the complementarity determining region 3 (CDR3) hypervariable region of immunoglobulin heavy chain (VH-CDR3) alone or in combination with at least another immunomodulating sequence. The DNA sequences are useful as therapeutic vaccines for thronic B-lymphoproliferative disorders in mammals, preferably humans. A recombinant plasmid expression vector containing a DNA sequence of the invention is useful as a therapeutic vaccine or for the manufacture of a vaccine effective against chronic B-lymphoproliferative disorders expressing the surface idiotypic immunoglobulin on B-cells in mammals, preferably humans. An efficient, safe and easily reproducible DNA-based immune response against B-lymphoproliferative pathologies can be achieved. The present sequence represents a PCR primer used to amplify thumans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of DNA sequences coding for hypervariable region (VH- complementarity determining region 3 (CDR3)) of idiotypic immunoglobulin expressed on B-cells of chronic B- lymphoproliferative disorders, as therapeutic
                                                                                                                                                                                                                                                               Human; immune response; chronic B-lymphoproliferative disorder; CDR: complementarity determining region 3; hypervariable region; B-cell; immunoglobulin heavy chain; VH-CDR3; idiotypic immunoglobulin;
                                                                                                                                                                                                                        Human heavy chain CDR3 variable region, PCR primer VH1-3-5-7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 15.4; DB 1; Length 21; 76.2%; Pred. No. 1.5e+03; rative 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 3 A; 3 C; 8 G; 5 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Fig 1B; 30pp; English.
                                                                                      ABK89992 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-2001; 2001WO-IT000014.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JAN-2001; 2001WO-IT000014.
                                                                                                                                                                                                                                                                                                                                        cytostatic; PCR; primer; ss.
                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fazio VM, Saglio G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-583654/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (FAZI/) FAZIO V M. (SAGL/) SAGLIO G.
                                                                                                                                                                                                                                                                                                                                                                                                                               WO200255559-A1
                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                             21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-JUL-2002.
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                                                                                                                                 ABK89992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              vaccine.
                                          RESULT 1253
                                                                 4BK89992
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The invention relates to an isolated polymucleotide comprising a sequence having 2618 bp encoding a phosphatidic acid-preferring phospholipase Al colympetide comprising a sequence having 872 amino acids or which is at least 88% identical to it; or a fragment, or a variant of the colymucleotide. The pharmaceutical composition comprising the expression vector or the reagent is useful for preparing a medicament for modulating the activity of a phosphatidic acid-preferring phospholipase Al protein in a disease such as CNS, genitourinary, cardiovascular or haematological disorder, asthma, cancer or chronic obstructive pulmonary disease. The colymucleotide of the invention can also be used in dispnostic assays for detecting diseases or abnormalities, or susceptibility to diseases related to the presence of mutations in the nucleic acid sequences that encode the enzyme. The polymucleotide of the invention can also be used in gene therapy to treat such disorders. This polymucleotide sequence represents a PCR primer of the phosphatidic acid-preferring phospholipase Al protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, multidrug resistance-associated protein 1; MRP-1; ss; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.4; DB 1; Length 21; Pred. No. 1.5e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 5 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human MRP-1 polymorphic DNA region #276.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2363 GIGCCIGIGCGIGCG 2379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-NOV-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1255
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Gaps

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853 GAGGAGGAGCTGGTGGAGGCT 873 SAGGTGCAGCTGGTGSAGTCT 21

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16; Conservative

Local Similarity

Best Loc Matches

AAL41510 standard; DNA; 21 BP

RESULT 1254 AAL41510 ID AAL41510 XX

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Example 9; Page 84; 198pp; English
                                                                                                                                                                                                                                                                                                3 GGATGGCACAGGGCTGGTG 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human NOVX probe SEQ ID NO:265.
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2001US-0349575P.
2001US-0346357P.
2002US-0373260P.
                                                                                                                                                                                                                                                           1 Similarity 84.2%;
16; Conservative
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2001US-0327449P.
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2001US-0330142P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002WO-US031373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001US-0327917P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-0339266P
                                                                                                                                                                                                                                                                                                                                                                                ADA05905 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
 WPI; 2002-657475/70.
                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003029424-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-OCT-2002;
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09-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-OCT-2001;
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17-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                            06-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                       ADA05905;
                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                        RESULT 1257
ADA05905/c
                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                           1) polynucleotide. The polynucleotide is useful in an in vitro method for identifying a single nucleotide polymorphism and for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of MRP-1 or for identifying and obtaining an inhibitor of the activity of a molecular variant of MRP-1. The sequences are useful for disapnosing a disorder related to the presence of a molecular variant of MRP-1 or susceptibility to such a disorder, where the disorder is cancer (particularly renal cancer) or a disease related to multidrug
                                                                                                                                                                                                             Novel multidrug resistance-associated protein 1 polynucleotide useful for diagnosis and treatment of cancer and multidrug resistance related diseases, and for identifying single nucleotide polymorphisms.
                                                                                                                                                                                                                                                                                   The invention relates to a multidrug resistance-associated protein 1 (MRP
                                                                                                                                                                                                                                                                                                                                                                                          resistance. This sequence represents a human MRP-1 polymorphic DNA region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; multidrug resistance-associated protein 1; MRP-1; ss; cancer; renal cancer; cytostatic; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.4; DB 1; Length 21;
44.2%; Pred. No. 1.5e+03;
ve 1; Mismatches 2; Indels
 renal cancer; cytostatic; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 21 BP; 5 A; 10 C; 3 G; 2 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brinkmann U, Hoffmeyer S, Mornhinweg E;
                                                                                                                                                                 Brinkmann U, Hoffmeyer S, Mornhinweg E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human MRP-1 polymorphic DNA region #275.
                                                                                                                                         (EPID-) EPIDAUROS BIOTECHNOLOGIES AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPID-) EPIDAUROS BIOTECHNOLOGIES AG.
                                                                                                                                                                                                                                                           Example 9; Page 84; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 GGATGGCACAGGGCTGGTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 GGGTGGCACRGTGCTGGTG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS67013 standard; DNA; 21 BP.
                                                                                            25-JAN-2002; 2002WO-EP000796.
                                                                                                                  26-JAN-2001; 2001EP-00101651
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                        WPI; 2002-657475/70.
                                              WO200259142-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200259142-A2
                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-AUG-2002.
                                                                      01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS67013;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1256
ABS67013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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The invention relates to a multidrug resistance-associated protein 1 (MRP of invention relates to a multidrug resistance-associated protein 1 (MRP for identifying as ingle nucleotide polymorphism and for identifying and obtaining a pro-drug or drug capable objection and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of MRP-1 or for identifying and obtaining an inhibitor of the activity of a molecular variant of MRP-1. The sequences are useful for diagnosing a disorder related to the presence of a molecular variant of MRP-1 or susceptibility to such a disorder, where the disorder is cancer (particularly renal canner) or a disease related to multidrug resistance. This sequence represents a human MRP-1 polymorphic DNA region
Novel multidrug resistance-associated protein 1 polynucleotide useful for diagnosis and treatment of cancer and multidrug resistance related diseases, and for identifying single nucleotide polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunomodulator; cytostatic; nootropic; neuroprotective;
antiparkinsonian; antilipaemic; gene therapy; human disease;
metabolic disorder; diabetes; obesity; infection; cachexia; cancer;
neurodegenerative disorder; Alzheimer's disease; Parkinson's disease;
immune disorder; haematopoietic disorder; dyslipidaemia; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.4; DB 1; Length 21;
4.2%; Pred. No. 1.5e+03;
ve 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 2 A; 3 C; 10 G; 5 T; 0 U; 1 Other;
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New NOVX polypeptides and nucleic acids, useful for diagnosing, preventing or treating NOVX-associated disorders, e.g. diabetes, obesity, cancer or dyslipidemia, and in chromosome mapping, tissue typing or
                                                                                                                                                            Smithson G, Millet I, Peyman JA, Kekuda R, Ju J, Li L, Guo X;
Patturajan M, Spytek KA, Edinger SR, Ellerman K, Malyankar UM;
Ort T, Gorman L, Zerhusen BD, Anderson DW, Zhong M, Catterton E;
Ji W, Miller CE, Rastlelli L, Stone DJ, Pena CEA, Shenoy SG;
Shinkets RA, Rothenberg ME, Leach MD, Agee ML, Berghs C, Dipippo VA;
Eisen AJ, Gangolli EA, Rieger DK, Spaderna SK;
                            19-APR-2002; 2002US-0373884P.
22-APR-2002; 2002US-0374977P.
16-MAY-2002; 2002US-0381037P.
16-MAY-2002; 2002US-0381042P.
17-MAY-2002; 2002US-0381042P.
                                                                                       28-MAY-2002; 2002US-0383656P.
29-MAY-2002; 2002US-038381P.
25-UTNA-2002; 2002US-031335P.
01-OCT-2002; 2002US-00262511.
                    2002US-0373826P
                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                      WPI; 2003-381626/36.
                                                                                                                                                                                                                                                                                        pharmacogenomics.
19-APR-2002; 2
19-APR-2002; 2
19-APR-2002; 3
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The present invention describes NOVX proceins, where X can be 1 to 55

(e.g. NOV1). Also described: (1) a composition comprising a polypeptide described above and a carrier; (2) a kit comprising, in one or more containers, the composition described above; (3) an isolated nucleic acid molecule davoe; (3) an isolated nucleic acid molecule described above; (5) a cell comprising the nucleic acid molecule described above; (5) a cell comprising the nucleic acid molecule described above; (5) a cell comprising the polypeptide described above; (7) methods for determining the presence or amount of the above polypeptide or nucleic acid molecule in a disease associated with altered levels of expression of the above colypeptide or nucleic acid molecule in a first mammalian subject; (9) a method for identifying a potential therapeutic agent for use in treating a pathology that is related to an aberrant expression or above; (10) a method for identifying a potential therapeutic agent for use in treating a pathology that is related to an aberrant expression or abortwity of the polypeptide; (11) a method of secribed above; (12) methods of the above polypeptide or modulator of activity or of latency or predisposition to a pathology associated with the above polypeptide in a method for modulator of activity or of latency or predisposition to capathology associated with the above polypeptide in a manual, and (14) a method for producing the above polypeptide in a contenting a method for producing the above polypeptide in a contenting a method for producing the above polypeptide in a contention and antilipaemic activities, and can be used in gene therapy. The polypeptide is useful in manufacturing a medicament for treating a caid molecule may be used to diagnose, treat or prevent metabolic caid molecule may be used to diagnose, treat or prevent metabolic caid molecule may be used to diagnose and present and antilipaemic activities, and can be used to present and activity of activities, and can be used in gene therapy. The call conten disease, immune disorders, haematopoietic disorders and various dyslipidaemias. The nucleic acids can also be used as hybridisation probes in chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The present sequence represents a probe for a human NOVX sequence, which is used in an example from the present invention. Example C; Page 368; 586pp; English.

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                                         Gaps
                                         ö
0.4%; Score 15.4; DB 1; Length 21; 94.1%; Pred. No. 1.5e+03; ive 0; Mismatches 1; Indels
   Query Match 0.4
Best Local Similarity 94.1
Matches 16; Conservative
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Sequence 21 BP; 6 A; 10 C; 4 G; 1 T; 0 U; 0 Other;

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1815 TGGGTCCTGCTCTGGG 1831
                                            21 rececrecrecreses
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GlcNAc; sugar residue; glucosaminyl 6-O-sulfotransferase; 6-OST protein; heparan aulfate; 3-O-sulfated; 3-OST-u protein; antithrombin; blood flow; dialysis machine; thrombotic diasase; extracorporeal medical device; intracorporeal device; transplant; stent; prosthetic implant; hamster; Anticoagulant; ahrombolytic; sulfate; 6-0 position; N-acetylglucosamine; Hamster 6-OST-3 PCR primer target region #2. blood clotting; PCR; primer; ss AAL54133 standard; DNA; 21 28-MAR-2003 (first entry) AAL54133; RESULT 1258 AAL54133

WO200279258-A2.

10-OCT-2002

28-MAR-2002; 2002WO-US010172

30-AUG-2001; 2001US-0316289P. 28-MAR-2001; 2001US-0279523P

(MASI) MASSACHUSETTS INST TECHNOLOGY.

Beeler DL; Rosenberg RD, Zhang L,

WPI; 2003-129077/12.

Transferring sulfate to 6-0 position of N-acetylglucosamine sugar residue in a polysaccharide preparation, by contacting the preparation with glucosaminyl 6-0-sulfotransferase protein in presence of a sulfate donor.

Example 14; Page 29; 48pp; English.

The invention relates to a novel method for transferring a sulfate on to coloposition of a Nacetylglucosamine (GlcNAC) sugar residue in a polysaccharide preparation, comprising providing a polysaccharide conditions which pervised in the presence of a gluosaminyl 6-0-sulfotransferase (6-0ST) protein in the presence of a sulfate donor under conditions which permit 6-0ST protein to add a sulfate donor under conditions which permit 6-0ST protein to add a sulfate to the 6-0-position of a GlcNAC sugar residue. This method is cusful for enriching the portion of anticoagulant heparan sulfate present of and contacting the preparation with 6-0ST protein. Buriching the preparation with 6-0ST protein. Buriching the contacting the preparation with 6-0ST protein. Buriching the antithrombin-binding fraction of a heparan sulfate pool is useful in the production of anticoagulant heparan sulfate which have clinical applications as therapeutics, e.g. as an agent to treat or prevent thrombotic disease and to maintain blood flow in medical devices, e.g. dialysis machines. The 6-0-sulfated preparations and the heparan sulfate to coat surfaces of extracorporeal medical devices or intracorporeal composed improved by administration an anticoagulant, e.g. thrombotic disease and coat surfaces of extracorporeal medical devices or intracorporeal copic devices (e.g. transplants, stents or other prosthetic implants) to reduce colocating on those surfaces. The mutant CHO cell hyper-produces anticoagulant-active heparan sulfate. This polynucleotide sequence

Seguence 21 BP; 5 A; 9 C; 3 G; 4 T; 0 U; 0 Other;

0.4%; Score 15.4; DB 1; Length 21; 94.1%; Pred. No. 1.5e+03; Query Match Best Local Similarity

Gaps

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Indels

1;

0; Mismatches

Matches 16; Conservative

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Gaps

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ACC69489 to ACC69495 encode the human dual-specificity phosphatases
designated DSP-18a to DSP-19f and prototypical DSP-18pr given in ABR43450
CC cho ABR43456. DSP-18 proteins have the ability to dephosphorylate an
activated mitogen activated protein (MAP)-kinase. DSP-18 sequences have
cytostatic, immunosuppressive and antiallergic activities, and can be
cytostatic in be used for identifying antibodies and other modulators
cyparticularly inhibitors) of DSP-18 activity. The DSP-18 proteins may be
cyparticularly inhibitors of DSP-18 activity. The DSP-18 proteins are especially useful
conversally or treat diseases associated with cell proliferation of
coll, differentiation or survival. The DSP-18 substrates. A modulator of
coll, differentiation of a cell or modulating a proliferative response in a
cell, differentiation of a cell or survival of a cell; or for treating a
cell, differentiation of a cell or survival of a cell; or for treating a
cell, differentiation of a cell or survival of a cell; or for treating a
cell, differentiation of a cell or survival of a cell; or for treating a
cell, differentiation of a cell or survival of a cell; or for treating a
cell, differentiation of a cell or survival of a cell; or for treating
cell, differentiation of a cell or survival of a cell; or for treating
cell, differentiation of a cell or survival of a cell; or for treating
cell, differentiation of a cell growth, abnormal cell proliferation, or
cell cycle abnormalities) associated with DSP-18 activity. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; dual-specificity phosphatase; DSP-18; enzyme; cytostatic; immunosuppressive, antiallergic; MAP-kinase modulator; dephosphorylation; signal transduction modulator; cell prolliferation; cell differentiation; cell survival; prolliferative response; Duchenne muscular dystrophy; cancer; graft-versus-host disease; autoimmune disease; allergy; metabolic disease; abnormal cell growth; abnormal cell prolliferation; cell cycle abnormality; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New DSP-18 dual-specificity phosphatases, useful for modulating cell proliferation, differentiation or survival, or for identifying modulators of DSP-18 activity for treating e.g. cancer or graft-versus-host disease
    ·,
    Indels
    1;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human DSP-18 PCR primer SEQ ID NO:26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 59; 113pp; English.
                                                                           TCACCTTCAAGGACCTG 1557
                                                                                                                                                                                                                                                                                                                                                                      ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-MAY-2001; 2001US-0291476P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-MAY-2002; 2002WO-US015906
                                                                                                                                                                                                                                                                                                                                                                      ACC69504 standard; cDNA; 21
                                                                                                                                                                TCACCCTCAAGGACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-371819/35.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wei B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003025196-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a patient.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Luche RM,
                                                                                1541
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACC69504;
                                                                                                                                                                N
                                                                                                                                                                                                                                                                                      RESULT 1259
                                                                                                                                                                                                                                                                                                                        ACC69504/C
ID ACC69504/C
XX
AC ACC6950
XX
XX
B Human |
XX
Hom os |
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    Matches
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Length 21;

Score 15.4; DB 1; Pred. No. 1.5e+03;

0.4%; 94.1%;

Query Match Best Local Similarity

Seguence 21 BP; 5 A; 3 C; 8 G; 5 T; 0 U; 0 Other;

example from the present invention

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The present invention describes an antibody expression library derived from a patient that has been immunochallenged with one or more foreign antigens associated with a particular disease or foreign agent. Also described: (1) a method for producing the expression library derived from the patient as described above; (2) libraries of cloned nucleic acid fragments comprising the variable heavy chain regions and/or variable cight fragments comprising the variable heavy chain regions and/or variable cight fragments, of antibody spense derived from the antibody-producing cells of immunochallenged patients; (3) a method or isolating from an antibody expression library one or more antibody molecules that is a specific binding patrent for a carget antigen; (4) a method of manufacturing a specific antibody comparison of target antibody molecule identified, manufactured or formulated as above; (6) an enthod of traating, diagnosing or imaging a patient; (7) a method to develop and/or produce a recombinant vaccine; (8) a comprising one or more dummy nucleic acid fragments located in the parts of the vector into which the nucleic acid inserts encoding a desired of that they are not in reading frame with the other parts of the expression to polypeptide will be cloned, where the dummy fragments are positioned such that they are not in reading frame with the other parts of the expression conducts. The antibody expression library has cytostatic, immunosuppressive and antiallergic activities, and can be used in gene products. The antibody expression florary and can be used in gene conducts. The antibody expression florary with different target antigens to identify cross-reactive antibodies, for developing vaccines and identifying the immunosens of a vaccine that stimulate the most
                                                                                                                                                                                                                                                                                                                     antibody expression library; antibody; vaccine; cytostatic; immunosuppressive; antiallergic; gene therapy; cancer; autoimmune disease; allergic disease; target antigen; PCR primer; V-gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antibody expression library derived from a patient that has been immunochallenged with one or more foreign antigens, useful for treating cancer, or autoimmune and allergic disease.
                                                                                                                                                                                                                                                                               V-gene primary amplification PCR primer N5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kausmally L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 64; 86pp; English.
1880 AGCICTICAAGCIGCIG 1896
                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-MAY-2002; 2002GB-00011015.
29-NOV-2002; 2002GB-00027977.
06-MAR-2003; 2003US-00379996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-MAY-2003; 2003WO-GB002073.
                                                                                                                                                  ADG77085 standard; DNA; 21
                                        18 AACTCTTCAAGCTGCTG
                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brekke OH, Stacy J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-012090/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AFFITECH AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003095491-A2.
                                                                                                                                                                                                                                        11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                           ADG77085;
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(OWEN/)
                                                                                                          RESULT 1260
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cancers including breast, retinoblastoma, Wilm's tumour and lymphoma. Accordingly, these compositions exhibit cytostatic, antidiabetic, antidiabetic, antiarflammatory, antigboriatic, antirheumatic and antiarthritic activities. This oligonucleotide is a human Flk-1/KDR DNA oligo, a target for siRNA inhibition of the invention.

888888888

rheumatoid arthritis. Furthermore, it can be used to treat various

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Gaps

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1; Indels

0; Mismatches

94.18; 0.4%;

Query Match
Best Local Similarity 94.1
Matches 16; Conservative

3013

2997 CACCGCAGTTTTGTTTT

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Score 15.4; DB 1; Length 21; Pred. No. 1.5e+03;

Sequence 21 BP; 10 A; 2 C; 4 G; 5 T; 0 U; 0 Other;

vivlemore401-10.rng

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                                                                                                                                                                                                                                                                                                                                                                                                   human; ss; short interfering RNA; siRNA; angiogenesis; vascular endothelial growth factor; VEGF; VEGF receptor; Flt-1; Flk-1/KDR; kinase domain region; diabetic retinopathy; age-related macular degeneration; inflammatory disease; psoriasis; rheumatoid arthritis; cancer; breast; retinoblastoma; Wilm's tumour; lymphoma; cytostatic; antidiabetic; ophthalmological; antiinflammatory; antipsoriatic; antirheumatic; antiarthritic.
        identifying, selecting or manufacturing one or more antibody molecules that bind specifically to one or more target antigens. The expression vector is useful for producing an antibody expression library. The present sequence represents a PCR primar for the primary amplification V-gene, which is used in an example from the present invention for the generation and screening of meningococcal B vaccine libraries.
                                                                                                                                                                                                                                                                                                                                                                          Human Flk-1/KDR DNA sequence, a target for siRNA inhibition SeqID 532
                                                                                                                                                                 Gaps
effective and protective antibodies, and for detecting, isolating,
                                                                                                                                                                 ö
                                                                                                                                   0.4%; Score 15.4; DB 1; Length 21; 76.2%; Pred. No. 1.5e+03;
                                                                                                                                                                 3; Indels
                                                                                                          Sequence 21 BP; 3 A; 3 C; 9 G; 3 T; 0 U; 3 Other;
                                                                                                                                                                2; Mismatches
                                                                                                                                                                                          853 GAGGAGGAGCTGGTGGAGGCT 873
                                                                                                                                                                                                             1 GAGGTGCAGCTGKTGGAGWCY 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUL-2003; 2003WO-US022444.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JUL-2002; 2002US-0398417P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-NOV-2002; 2002US-00294228
                                                                                                                                                                                                                                                                                           ADJ97759 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                 06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                 16; Conservative
                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004009769-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                      ADJ97759;
                                                                                                                                      Query Match
                                                                                                                                                                 Matches
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This invention relates to novel compositions that comprise short interfering RNA (siRNA) molecules, which can be used to inhibit amagiogenesis. Specifically, it refers to siRNAs that target and cause RNAi-induced degradation of mRNA from human vascular endothelial growth factor (VEGF), the VEGF receptor (FIt-1) and the FIk-1/KDR (kinase domain region) genes, as well as mutants derived thereof. The present invention bind to the target mRNA, such that expression is inhibited and the target degraded. As such, siRNA administered in combination with a therapeutic agent is useful for treating diseases associated with angiogenesis and the overexpression of VEGF, which include diabetic retinopathy, agerelated macular degeneration, inflammatory disease, psortasis and rheumatoid arthritis. Furthermore, it can be used to treat various
                                                                                                                                                                                                                                                   human; 88; short interfering RNA; siRNA; angiogenesis; vascular endothelial growth factor; VEGF; VEGF receptor; Flt-1; Flk-1/KDR; kinase domain region; diabetic retinopathy; sege-related macular degeneration; inflammatory disease; psoriasis; rheumatoid arthritis; cancer; breast; retinoblastoma; Wilm's tumour; lymphoma; cytostatic; antidiabetic; ophthalmological; antiinflammatory; antipsoriatic; antirheumatic; antiarthritic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel short interfering RNA (siRNA) comprises sense and antisense RNA strands, useful for inhibiting expression of human vascular endothelial growth factor mRNA, for treating angiogenic disease, e.g. diabetic
                                                                                                                                                                                                                Human Flk-1/KDR DNA sequence, a target for siRNA inhibition SeqID 533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 533; 218pp; English.
                                                                                                  BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JUL-2002; 2002US-0398417P.
14-NOV-2002; 2002US-00294228.
18 CACCACAGTTTTGTTT
                                                                                                  ADJ97760 standard; DNA; 21
                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tolentino MJ, Reich SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 retinopathy and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-203472/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2004009769-A2
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                               06-MAY-2004
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                                                                                                                                         ADJ97760;
                                                           RESULT 1262
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This invention relates to novel compositions that comprise short interfering RNA (siRNA) molecules, which can be used to inhibit angiogenesis. Specifically, it refers to siRNAs that target and cause RNAsi-induced degradation of mRNA from human vascular endothelial growth factor (VEGP), the VEGF receptor (Flt-1) and the Flk-1/KDR (kinase domain describes sense and antisense RNA strands thereof. The present invention bind to the target mRNA, such that expression is inhibited and the target degraded. As such, siRNA administered in combination with a therapeutic agent is useful for treating diseases associated with angiogenesis and the overexpression of VEGF, which include diabetic retinopathy, agerrelated macular degeneration, inflammatory disease, psoriasis and

Novel short interfering RNA (siRNA) comprises sense and antisense RNA strands, useful for inhibiting expression of human vascular endothelial growth factor mRNA, for treating angiogenic disease, e.g. diabetic

Tolentino MJ, Reich SJ;

WPI; 2004-203472/19.

Disclosure; SEQ ID NO 532; 218pp; English.

retinopathy and cancer.

Gaps ;

Matches

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Accordingly, these compositions exhibit cytostatic, antidiabetic, ophthalmological, antinflammatory, antipsoriatic, antirheumatic and antiarthritic activities. This oligonuclectide is a human Flk-1/KDR DNA oligo, a target for siRNA inhibition of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        infectious disease; anorexia; cancer; cancer-associated cachexia; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; immune disorder; haematopoietic disorder; dyslipidaemia; metabolic syndrome X; wasting disorder; probe.
                                                                                                                                                        Score 15.4; DB 1; Length 21;
Pred. No. 1.5e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; human; NOVX; metabolic disorder; diabetes; obesity;
                                                                                                                  4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                             3013
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2001US-0328029P.
2001US-0328044P.
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2001US-0328849P.
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2002US-0391335P.
                                                                                                                                                        Query Match 0.4%;
Best Local Similarity 94.1%;
Matches 16; Conservative
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2002US-0373260P
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                                                                                                               Sequence 21 BP; 11 A; 3 C;
                                                                                                                                                                                                                                             2997 CACCGCAGTTTTGTTTT
                                                                                                                                                                                                                                                                                     21 CACCACAGITITIGITIT
                                                                                                                                                                                                                                                                                                                                                                                              ADN63068 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human NOVX probe #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2004038223-A1.
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09-OCT-2001;
09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-0CT-2001;
12-0CT-2001;
15-0CT-2001;
17-0CT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-OCT-2001;
29-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        ADN63068;
                                                                                                                                                                                                                                                                                                                                                      RESULT 1264
                                                                                                                                                                                                                                                                                                                                                                            ADN63068,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to novel compositions that comprise short interfering RNA (sirRA) molecules, which can be used to inhibit angiogenesis. Specifically, it refers to sirNAs that target and cause RNAsi-induced degradation of mRNA from human vascular endothelial growth factor (VBGP), the VBGP receptor (File-1) and the File-1/KDR (Kinase domain region) genes, as well as mutants derived thereof. The present invention describes sense and antisense RNA strands that form an RNA duplex and bind to the target mRNA, such that expression is inhibited and the target of degraded. As such, siRNA administered in combination with a therapeutic agent is useful for treating diseases associated with angiogenesis and the overexpression of VBGP, which include diabetic retinopathy, agerelated macular degeneration, inflammantory disease, psoriasis and rehematorid arthritis. Furthermore, it can be used to treat various cancers including breast, retinoblastoma, Wilm's tumour and lymphoma.
                                                                                                                                                                                                                        ö
cancers including breast, retinoblastoma, Wilm's tumour and lymphoma. Accordingly, these compositions exhibit cytostatic, antidiabetic, ophthalmological, antinflammatory, antipsoriatic, antirheumatic and antiarthritic activities. This oligonuclectide is a human Flk-1/KDR DNA oligo, a target for siRNA inhibition of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; 88; short interfering RNA; siRNA; angiogenesis; vascular endothelial growth factor; VEGF, VEGF receptor; Flt-1; Flk-1/KDR; kinase domain region; diabetic retinopathy; age-related macular degeneration; inflammatory disease; psoriasis; rapenmatoid arthritis; cancer; breast; retinoblastoma; Wilm's tumour; lymphoma; cytostatic; antidiabetic; ophthalmological; antiinflammatory; antipsoriatic; antiarthritic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel short interfering RNA (siRNA) comprises sense and antisense RNA strands, useful for inhibiting expression of human vascular endothelial growth factor mRNA, for treating angiogenic disease, e.g. diabetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Flk-1/KDR DNA sequence, a target for siRNA inhibition SeqID 531
                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                         0.4%; Score 15.4; DB 1; Length 21; 94.1%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                      1; Indels
                                                                                                                                 Sequence 21 BP; 9 A; 3 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 531; 218pp; English.
                                                                                                                                                                                                                                                              2997 CACCGCAGTTTTGTTT 3013
                                                                                                                                                                                                                                                                                                                                                                                                                  BP
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                                                                                                                                                                                              94.18;
                                                                                                                                                                                                                                                                                                                                                                                                                ADJ97758 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Folentino MJ, Reich SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             retinopathy and cancer.
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                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-MAY-2004
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                                                                                                                                                                                                                                                                                                       17
                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ97758;
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MILLET I. PEYMAN J A. SMITHSON G.

(MILL/) (/LIWS

represents a human NOVX probe.

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The invention relates to isolated NoVX polypeptides and polymucleotides.

NoVX polypeptides and polymucleotides are used to prevent, diagnose or treat a medical condition in human related to the aberrant expression and activity of NoVX polypeptides. For example, NoVX polypeptides and activity of NoVX by supplementing the patient our polymucleotides may be used to treat disorders associated with decreased compression or activity of NoVX by supplementing the patient our production or to rectify mutations. Conversely, antisense NA molecules may be administered to down regulate expression of NoVX polypeptides by blodymucleotides and complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar sequences in samples, and so which patients may be in need of restorative therapy. NoVX polypeptides may also be used as antigens in the production of antibodies and in assays to identify modulators anti-NoVX polypeptides and antagonists of the expression and activity of NoVX. The anti-NoVX polypeptides and antagonists and antagonists and antagonists of the expression and activity of NoVX canti-NoVX polypeptides antibodies and antagonists may also be used as diagnostic agents for detecting the presence of NoVX in samples. NoVX colypeptides and polymucleotide expression and activity of NoVX colypeptides and polymucleotide may be used in this way to prevent. Colisease, ancrexia, cancer associated cachexia, metabolic disorders, harbaimer's Disease, Parkinson's Disorders, and the various dyslipidaemias, metabolic disorders, has matabolic syndrome X and colisturbances associated with obesity, the metabolic syndrome X and washing disorders associated with obesity, the metabolic syndrome X and washing disorders associated with obesity, the metabolic syndrome X and washing disorders associated with obesity, the metabolic syndrome X and washing disorders associated with obesity, the metabolic syndrome X and context.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated NOVX polypeptides and nucleic acids, useful for preventing, diagnosing and treating e.g. cancer, diabetes and Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                            STONE D J.
PENA C E A.
SHENOY S G.
SHIMKETS R A.
ROTHENBERG M E.
                                                                   PATTURAJAN M.
SPYTEK K A.
EDINGER S R.
                                                                                                                          ELLERMAN K.
MALYANKAR U M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EISEN A.
GANGOLLI E A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIEGER D K.
SPADERNA S K.
                                                                                                                                                                                                                                                                                                                                                                                                                                           AGEE M L.
BERGHS C.
DIPIPPO V A.
                                                                                                                                                                                                 ZERHUSEN B D.
ANDERSON D W.
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                                                                                                                                                                                                                                                       CATTERTON E.
                                                                                                                                                                                                                                                                                         MILLER C E.
                                                                                                                                                                                                                                                                                                              RASTELLI L
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                                                                                                                                                                                GORMAN L.
KEKUDA R.
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                                                     GUO X.
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(RIEG/)
(SPAD/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated, mutant murine or human phospholipase C (PLC) gamma-2 protein. The mutant PLCgamma-2 proteins and nucleic acids that encode them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate PLCgamma-2 expression. For example, the peptides and nucleic acids may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of PLCgamma-2 by expressing inactive proteins or to supplement the patients administered to down regulate PLCgamma-2. Conversely inhibitory nucleic acids may be administered to down regulate PLCgamma-2 expression. The PLCgamma-2 proteins and nucleic acids may also be used in assays to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modulators of PLCgamma-2 expression and activity. The antibodies and antagonists may also be used to down regulate expression and activity. Disorders that may be prevented, diagnosed and/or treated by the above methods include, e.g. Wegener's granulomatosis, Churg-Strauss syndrome or Polyarteritis nodosa. This sequence corresponds to a PCR primer to amplify the mouse phospholipase C gamma 2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mutant murine or human Phospholipase C (PLC) gamma-2 proteins, useful for preventing, diagnosing and treating inflammatory diseases, e.g. Wegener's granulomatosis, Churg-Strauss syndrome or Polyarteritis nodosa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; primer; anti-inflammatory; dermatological; osteopathic; ophthalmological; antiulcer; murine; human; phospholipase C gamma-2 protein; diagnosis; Wegener's granulomatosis; Churg-Strauss syndrome; Polyarteritis nodosa.
                                                                                                                      Gaps
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                                                                              Score 15.4; DB 1; Length 21;
Pred. No. 1.5e+03;
                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schroeder A, Yu P, Hanke P;
                                        Sequence 21 BP; 6 A; 10 C; 4 G; 1 T; 0 U; 0 Other;
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                                                                                                                    0; Mismatches
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                                                                                                                                                              1815 TGGGGTCCTGCTCTGGG 1831
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                                                                              Query Match
Best Local Similarity 94.1%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                       ADO80395 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                     29-JUL-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-449493/42.
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004020619-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus.
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Dipippo VA;

vivlemore401-10.rng

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Identifying extendible primers for use in identification, or classification of a nucleic acid of an organism, allele or gene such as class 1/2 HLA comprises identifying all possible nucleotide sequences of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides a method for identifying a set of extendible primers which can be used in the identification, typing and classification of genes. This can then be used to predict protein sequence and structure, in organ donation to match the organ with the receiver, and to identify bacteria in a sample. The method can be used type the human leukocyte antigen genes (HLA) and 16s rRNA genes in
                                                                       DNA sequence analysis, sequencing, protein sequence, protein structurgene typing, organ donation, bacteria identification, 16s rRNA, HLA, human leukocyte antigen, PCR primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 25 BP; 2 A; 4 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               (AMSH ) AMERSHAM PHARMACIA BIOTECH AB.
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                        HLA DPB1 gene PCR primer #21.
                                                                                                                                                                                                                                                                                                                                            20-APR-2000; 2000WO-EP003636.
                                                                                                                                                                                                                                                                                                                                                                                              99EP-00303215.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ulfendahl P, Wong K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-679677/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific length.
                                                                                                                                                                                                                                      WO200065088-A2.
                                                                                                                                                                                                                                                                                                                                                                                              26-APR-1999;
                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an immunoglobulin molecule comprising of one or more heavy chain framework regions comprising HFR1, HFR2, HFR3 or HFR4 and one or more light chain framework regions comprising LFR1, LFR2, LFR3 or LFR4, and complementarity determining regions (CDRs) comprising CDR-H1, CDR-H2, CDR-H3 and/or CDRL1, CDR-L2 or CDR-L3. The immunoglobulin is useful in therapeutic or diagnostic assays comprising ELISA, phage display, tumour imaging or protein chips assay. Further, the immunoglobulin is useful in screening assays for detecting molecules that bind to the immunoglobulin molecule. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                    in molecule; heavy chain framework region; HFR1; HFR2; HFR3; imaging; protein chips assay; light chain framework region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New immunoglobulin molecule, useful in therapeutic or diagnostic assays comprising ELISA, phage display, tumor imaging or protein chips assay or in screening assays for detecting molecules that bind to the immunoglobulin molecule.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 22 BP; 4 A; 6 C; 9 G; 3 T; 0 U; 0 Other;
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                        TGTCCTGTGCCTACCAG 1575
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                                                                                                                                                                                                                                                                                                                                                                                                                    immunoglobulin molecule; heavy
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                                                                 TGTCTTGTGCCTACCAG
                                                                                                                                                                                                        ADI20640 standard; DNA; 22
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                                                                                                                                                                                                                                                                                                                                                                  Primer of the invention #1
                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.4
Best Local Similarity 94.1
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schillberg S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-371805/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LFR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003025124-A2.
                                                                                                                                                                                                                                                                                                              22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhang MY,
Fischer R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                     1559
                                                                          18
                                                                                                                                                                                                                                                            ADI20640;
                                                                                                                                                       RESULT 1266
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AC ADIZ
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BE Prim
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                                                                                                                                                                                                                                         DNA sequence analysis; sequencing; protein sequence; protein structure; gene typing; organ donation; bacteria identification; 16s rRNA; HLA; human leukocyte antigen; PCR primer; ss.
                           Gaps
                           ..
0
Score 15.4; DB 1; Length 25;
Pred. No. 1.8e+03;
0; Mismatches 6; Indels
                                                 3264 TITIATITGCTTTGTCCTTTTCAG 3288
                                                                          1 rrrrrrrrrrrrrrrsccrrrrccag 25
                                                                                                                                        ВР
                                                                                                                                                                                                                   HLA DPB1 gene PCR primer #99.
                                                                                                                                                                                                                                                                                                                                                                                            99EP-00303215.
                                                                                                                                                                                                                                                                                                                                                                    20-APR-2000; 2000WO-EP003636.
 Query Match
Best Local Similarity 76.0%;
Matches 19; Conservative
                                                                                                                                       AAC96367 standard; DNA; 25
                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                            26-APR-1999;
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AAC95686 standard; DNA;

RESULT 1267

AAC95686

AAC95686;

BXXXB

26-FEB-2001 (first entry)

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The present invention relates to an immunostimulatory composition, which comprises at least one oligonucleotide comprising both an RNA region and a DNA region. The composition is useful for enhancing an immune response or inducing cytokines. It can be used as a vaccine adjuvant and in treating diseases, including pathogenic infection, (non-)malignant trumours (e.g. cancers of the brain, lung, ovary, breast, prostate or colon, or carcinomas and sarcomas), autoimmune diseases or allergies (e.g. allergic rhinitis, hay fever or food allergies), Lyme disease, hepatitis, HIV or malaria. The composition is also useful for treating, preventing or ameliorating the symptoms resulting from exposure to a biowarfare agent, e.g. Boola, Anthrax or Listeria. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Triplex formation; DNA detection; triple helix; identification; bacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a polynucleotide that is able to form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Triple helix third strand of n-myc gene nucleotides 4791-4725.
                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.4; DB 1; Length 30; 76.0%; Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                    Sequence 30 BP; 26 A; 1 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                          3310 TTTTTCTTTAGGAGATTTATTTTT 3334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Col 13-14; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                               29 TITITITITIAGCTITITITI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PROF-) PROFILE DIAGNOSTIC SCI INC
                                  Example 1; Page 30; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93US-00173489.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX14633 Standard; DNA; 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-MAR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                        Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oncogene; virus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wang C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-130384/11.
                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
HIV infection.
                                                                                                                                                                                                                                                                               the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-DEC-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX14633,
                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1270
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                                                                                                                                                                                                                            The present invention provides a method for identifying a set of extendible primers which can be used in the identification, typing and classification of genes. This can then be used to predict protein sequence and structure, in organ donation to match the organ with the receiver, and to identify bacteria in a sample. The method can be used to type the human leukocyte antigen genes (HLA) and 16s rRNA genes in
                                                                                                   Identifying extendible primers for use in identification, or classification of a nucleic acid of an organism, allele or gene such as class 1/2 HIA comprises identifying all possible nucleotide sequences of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New immunostimulatory compositions comprising RNA/DNA hybrid oligonucleotides, useful for enhancing an immune response or inducing cytokines, particularly for treating diseases, e.g. cancer, allergy or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA/RNA hybrid; phosphorothioate backbone; immunostimulatory; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 infection; allergy; cancer; hypersensitivity; bio-warfare; immunostimulant; antiallergic; cytostatic; antimicrobial; anti-HIV; immunosuppressive; protozoacide; virucide; hepatotropic; gene therat antiinflammatory; antibacterial; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.4; DB 1; Length 25;
Pred. No. 1.8e+03;
0; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA/DNA hybrid assay control oligonucleotide SEQ ID NO: 11
                                                                                                                                                                                                                                                                                                                                                                       Sequence 25 BP; 2 A; 4 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3264 TTTTATTTGCTTTGTCCTTTTCAG 3288
(AMSH ) AMERSHAM PHARMACIA BIOTECH AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 rrrrrrrrrrrrrrrsccrrrrcag 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                           Claim 14; Page 50; 66pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Klinman DM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABL35105 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                        ch
1 Similarity 76.0%;
19; Conservative
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Matches 19; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-130570/17.
                                                                   WPI; 2000-679677/66.
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                                                                                                                                                        specific length.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified base
                                 Ulfendahl P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL35105;
                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Gaps

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Indels

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RNA template, C(UA)3G used to direct RNA synthesis by HCV RNA polymerase.
DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of
                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus; HCV replicase; non-structural protein 5B; NS5B;
                                                                                                                                                             .
                                                                                                                                   0.4%; Score 15.4; DB 1; Length 35; 76.0%; Pred. No. 2.3e+03; tive 0; Mismatches 6; Indels
                                                                                                              Sequence 35 BP; 0 A; 4 C; 1 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                     3262 TATITIATITGCTTIGICCTTTTC 3286
                                                                                                                                                                                                           rrrrrrrrrrrrrrrrrrrrrrrrrrrrr
                                                                                                                                                                                                                                                                                                                                                                                 lead compound; RNA polymerase; ss.
                                                                                       oncogenes and Hepatitis B virus
                                                                                                                                                                                                                                                                       AAD27122 standard; RNA; 36 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-00309670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-00309670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhong W, Hong Z, Lau JYN;
                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                           19; Conservative
                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-096587/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZHON/) ZHONG W.
(HONG/) HONG Z.
(LAUJ/) LAU J Y N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAY-1999;
                                                                                                                                                                                                                                                                                                                      09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                US6322966-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-NOV-2001.
                                                                                                                                                                                                                                                                                              AAD27122;
                                                                                                                                      Query Match
                                                                                                                                                                                                                                            Matches
  88888888888888
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The present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemloop at the 3' end capable of forming a copyback structure, a HCV non-structural protein 5B (NSSB), ATP, GTP, CTP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is and UTP nucleoside triphosphates (NTPS), where one of the NTP is and UTP nucleoside triphosphates to the identification of optimal compounds or replacements of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds correctise the biological relevance of lead compounds. The Correctise the biological relevance of lead compounds The COPT NASSB polymerase mechanistically and kinetically and for designing small RNA templates can be used for developing a system to characterise the co-crystatalise with HCV NSSB polymerase. The assay system of the invention is useful for detected by evaluating an sutoradiograph of reaction products separated by Gel electrophoresis. The present sequence is RNA template, CC(AU) 2GG used to direct RNA synthesis cused in the exemplification of the invention
                                                                                                                                                                                                            Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Fig 1C; 10pp; English.
Zhong W, Hong Z, Lau JYN;
                                                                                                              WPI; 2002-096587/13.
                                                                                                                                                                                                                                                                                                                                                                                                buffer.
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The present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemhloop at the 3' end capable of forming a copy-back structure, a HCV non-structural protein 5B (NSSB), AFP, GTP, TP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is radiolabelled and an assay buffer that supports replication activity of NSSB. The invention also relates to the identification of optimal properties of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds or to characterise the biological relevance of lead compounds. The optimal RNA templates can be used for developing a system to characterise HCV NSSB polymerase mechanistically and kinetically and for designing system of the invention is useful for detecting HCV replicase activity. The nucleic acid synthesised by NSSB is detected by evaluating an

Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates

Example 1; Fig 1C; 10pp; English.

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present sequence is RNA template, C(UA)3G used to direct RNA synthesis by RNA polymerase proteins of HCV, BVDV and polivirus. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                             RNA template CC(AU)2GG for directing RNA synthesis by HCV RNA polymerase.
                                                                                         Gaps
                                                                                                                                                                                                                                               Hepatitis C virus; HCV replicase; non-structural protein 5B; NS5B;
                                                                                         .
0
                                                                     Length 36;
                                                                    Query Match
0.4%; Score 15.4; DB 1; Length 3
Best Local Similarity 66.7%; Pred. No. 2.4e+03;
Matches 22; Conservative 0; Mismatches 11; Indels
                                                    Sequence 36 BP; 29 A; 1 C; 3 G; 0 T; 3 U; 0 Other;
                                                                                                          3302 CTATAGGATTTTTTTAGGAGATTTATTTTT 3334
                                                                                                                            CIAIAIAGITITITITITITITITITITIT 4
                                                                                                                                                                                                                                                        lead compound; RNA polymerase; ss.
                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                               99US-00309670.
                                                                                                                                                                                                                                                                                                                                                 99US-00309670.
                                                                                                                                                                        AAD27123 standard, RNA; 36
                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                          (HONG/) HONG Z.
(LAUJ/) LAU J Y N.
                                                                                                                                                                                                                                                                                                                                                                   (ZHON/) ZHONG W.
                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                               11-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                 11-MAY-1999;
                                                                                                                                                                                                                                                                                                             27-NOV-2001.
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                                                                                                                                                       RESULT 1272
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Gaps

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Indels

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No. 2.4e+03;

Pred. No. 2.46); Mismatches

76.0%;

Conservative

Local Similarity ses 19; Conserv

Best Loc Matches

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The present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemhoop at the 3' end capable of forming a copyback structure, a HCV non-structural protein 58 (NSSB), ATP, GTP, TP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is cradiclabelled and an assay buffer that supports replication activity of NSSB. The invention also relates to the identification of optimal properties of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds or to characterise the biological relevance of lead compounds. The optimal RNA templates can be used for developing a system to characterise (C small NA menlecules to co-crystallise with HCV NSSB polymerase mechanistically and kinetically and for designing system of the invention is useful for detecting HCV replicase activity. The nucleic acid synthesised by NSSB is detected by evaluating an incompleted.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     autoradiograph of reaction products separated by gel electrophoresis. The present sequence is RNA template, (AU)2 used to direct RNA synthesis by RNA polymerase proteins of HCV, BVDV and polivirus. This sequence is used
                                                                                                                                                                                                                                                                                                                                                        RNA template, (AU) 2 used to direct RNA synthesis by HCV RNA polymerase.
                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates.
                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis C virus, HCV replicase, non-structural protein 5B, NS5B, lead compound; RNA polymerase; ss.
                                                                       ö
                                      Length 36;
                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 37 BP; 33 A; 0 C; 2 G; 0 T; 2 U; 0 Other;
Sequence 36 BP; 28 A; 2 C; 4 G; 0 T; 2 U; 0 Other;
                                                                         11;
                                    DB 1;
                                                                                                            3302 CTATAGGATTTTTCTTTAGGAGATTTATTTTT 3334
                                  Score 15.4; DB 1;
Pred. No. 2.4e+03;
0; Mismatches 11
                                                                                                                                                36 ccararderrrrrrrrrrrrrrrrrrrrrrrrr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present sequence is RNA template, (AU) 2 RNA polymerase proteins of HCV, BVDV and in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Fig 2A; 10pp; English.
                                                                                                                                                                                                                                           AAD27125 standard; RNA; 37 BP.
                                      0.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-00309670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-00309670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lau JYN;
                                                                                                                                                                                                                                                                                                                     (first entry)
                                                        Best Local Similarity 66.7
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-096587/13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhong W, Hong Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZHON/) ZHONG W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HONG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JS6322966-B1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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                                                                                                                                                                                                                                                                                AAD27125;
                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (LAUJ/)
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                                                                                                                                                                                                         RESULT 1273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n using novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                            OCR; polymerase chain reaction; paternity; maternity; human; pedigree, inkage analysis; genetic disease; animal; plant; breeding; locus;
                                                                                                                                                                                                                                                                           repeat sequence; genetic marker; primer; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the marker clone Mdf111 which contains the repeat sequence having ocumula: TGAGACCCTG(AC)15.5. (Updated on 25-MAR-2003 to correct PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ch
1 Similarity 61.0%; Pred. No. 2.56+03;
25; Conservative 0; Mismatches 12. 7-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3047 TGGGCCCCTGGCACTCTTGTTCCCACACCCCAACACTTCCA 3087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 TGAGACCCTGACACACACACACACACACACACACA 41
                                                                                                                                                                                                                                           Repeat sequence from polymorphic marker clone Mfd111,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 41 BP; 18 A; 18 C; 3 G; 2 T; 0 U; 0 Other;
3261 ATATTTTATTTGCTTTGTCCTTTT 3285
                              Claim 1; Col 13-14; 186pp; English.
                                                                                                                                                                                                                                                                                                                                    hybridisation; chromosome; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            89US-00341562.
91US-00754351.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00222177.
                                                                                                                       AAT65782 standard; DNA; 41
                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                          (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-042299/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-APR-1989;
05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-APR-1994;
                                                                                                                                                                                        25-MAR-2003
17-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                       JS5582979-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                         10-DEC-1996.
                                                                                                                                                         AAT65782;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                     RESULT 1274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                       AAT65782
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0.4%; Score 15.4; DB 1; Length 37;

Query Match

RESULT 1275

AAT65783;

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Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer directed primer, comprising of a perfect cpd. simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor segment. The present sequence is an example of a SSR primer, which is flanked at its 5'-end by degenerate nucleotides. The method represents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified amplified fragment length polymorphism assay - for detection of
                               adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; 5'-anchored; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; 5'-anchored; ss.
Detection; polymorphism; perfect compound simple sequence repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 15.2; DB 1; Length 18;
33.8%; Pred. No. 1.4e+03;
ve 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5'-anchored simple sequence repeat primer VHV(GT)7.5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 0 A; 0 C; 7 G; 8 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphism esp. in micro:satellite regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 77; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (DUPO ) DU PONT DE NEMOURS & CO E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2332
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                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00346456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2317 CTGTGTGTGTGTGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vogel JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      germplasm comparisons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-277795/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            8-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                            21-NOV-1995;
                                                                                                                                                                                                                                  WO9617082-A2
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                                                                                                                                                                                                                                                                                                         06-JUN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-1996
                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT2791
MAKE WE WE WERE WELL WAS A STANFORD 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ઠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           repeats, especially for use in e.g paternity or maternity teeting, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAT65798-76647. Those clones where the repeat sequence has been determined are shown in AAT65704-797. This repeat sequence has the marker clone Mdf112 which contains the repeat sequence having the formula: CCACCCCCAG(CR)24.5. (Updated on 25-MAR-2003 to correct PP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection of polymorphic genetic markers of the form (dG-dA)n(dG-dT)n - veing novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                  Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3047 TGGGCCCCTGGCACTCTTGTTCCCACACCCCAACACTTCCA 3087
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                                                                                                                                                                                                                                                                      Repeat sequence from polymorphic marker clone Mfd112.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5'-anchored simple sequence repeat primer HVH(TG)7.5,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 41 BP; 18 A; 18 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Col 13-14; 186pp; English
                               AAT65783 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                hybridisation; chromosome; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  89US-00341562.
91US-00754351.
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Best Local Similarity 61.00,
                                                                                                                                                               (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-APR-1994;
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05-SEP-1991;
                                                                                                                                                               25-MAR-2003
17-JUN-1997
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RESULT 1276

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Weber JL;

AAT65783

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Gaps

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acid molecule encoding human Survivin, where the antisense coligonuclectide inhibits the expression of human Survivin. These coligonuclectides are used in the treatment of an animal suffering from a disease or condition associated with Survivin, e.g. a hyperproliferative condition such as cancer, and comprises administering a therapeuticially or prophylactically effective amount of the antisense oligonuclectides can also be used to treat a human suffering from a cligonuclectide so that expression of Survivin is inhibited. The coligonuclectides can also be used to treat a human suffering from a disease or condition characterised by a reduction in apoptosis comprising administering the antisense oligonuclectide to a human. In addition, the antisense oligonuclectide and a cytotoxic chemotherapeutic agent e.g. taxol or cisplatin, can be used to modulate apoptosis, cytokinesis or the cell cycle, or inhibit the proliferation in a cancer cell by contacting the cell with the antisense oligonuclectide. AAS21151-AAS21768 represent Survivin nucleic acids, and antisense oligonuclectides targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequences given in AAQ28303 and AAQ28309-13 are PCR primers which were used to amplify and isolate the Entomopoxvirus Amsacta moorei (AmEDV) spheroidin DNA sequence and associated regulatory regions. The isolated sequence contained six open reading frames encoding the spheroidin protein itself and also other structural or regulatory genes associated with spheroidin. BPV spheroidin has no significant amino acid homology to any previoualy reported protein. It is a non-essential protein which makes it desirable as a site for the insertion of exogenous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New viral vectors and chimeric vaccines - comprise entomopoxvirus expression system contg. spheroidin or thymidine kinase sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Entomopoxvirus; non-essential; regulatory sequences; vector; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 20 BP; 11 A; 2 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                             Survivin, used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymerase chain reaction; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3458 AAGTITATATATATATAT 3477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 ATGTTTATATATATATGT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amsacta moorei entomopoxvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AmEPV Spheroidin primer RM92.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYFL ) UNIV FLORIDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Moyer RW, Hall RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-316172/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9214818-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .9-FEB-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-SEP-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ28311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1279
       셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                     Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSN), and an adaptor directed primer, comprising a sequence complementary to an adaptor segment. The present sequence is an example of a SSR primer, which is flanked at its 5-end by degenerate nucleotides. The method represents modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and
                                                                                                                                                                                                  Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel antisense compounds for modulating the expression of Survivin and treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.2; DB 1; Length 18; 93.8%; Pred. No. 1.4e+03; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 0 A; 0 C; 8 G; 7 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse Survivin antisense oligonucleotide #57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Swayze EE,
                                                        (DUPO ) DU PONT DE NEMOURS & CO E I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 18; Page 62; 120pp; English
                                                                                                                                                                                                                                                                          Example 1; Page 77; 173pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2334 CGTGTGTGTGTGTG 2349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS21755 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001; 2001WO-US002939.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-FEB-2000; 2000US-00496694.
       94US-00346456.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bennett CF, Ackermann EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                      Morgante M, Vogel JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       germplasm comparisons
                                                                                                                                                    WPI; 1996-277795/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-488863/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200157059-A1
     28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS21755;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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RESULT 1278

AAS21755/ XX AAS2 XX AAS2 XX AAS2 XX MUS OS MUS OS SYNT XX WOOG XX XX CIS: XX

Matches

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The invention relates to antisense oligonucleotides targeted to a nucleic

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Gaps

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Indels

Gaps

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Indels

85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3;

1990 CCCACCTTCAAGCAGCTGGT 2009

17; Conservative

1 cccaccarrcaccacrecr 20

뮵

AAQ64159 standard; DNA; 20

Tyrosine kinase, receptor, proto-oncogene, trk, detection, diagnosis, antibody, treatment, tumour, antisense, ss.

Synthetic.

Primer for amplifying tyrosine kinase receptor coding sequence.

(first entry)

(revised)

25-MAR-2003 03-FEB-1995

AAQ64159;

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Best Local Similarity
                                                                                                                                                                                          RESULT 1281
                      Matches
                                                                                                                                                                                                                     AAQ64159/
                                                                                                                                                                                                                                                                                        임
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    임
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ò
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                      fragments of the surrounding DNA can be used as regulatory sequences if placed in operative association with foreign DNA. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The rpsL gene of M.leprae encodes the S12 protein of the small ribosomal subunit that is responsible for resistance to streptomycin. Two primers were designed based on this sequence and were used in a PCR amplification of M.tuberculosis DNA. Sequence analysis of the 306pp amplified fragment showed 28 differences between the rpsL gene from M.leprae (AAQ51533) and M.tuberculosis (AAQ61454). In streptomycin resistant strains, a single amino acid substitution due to a mutation in codon 43 (wild-type AAG mutated to AGG) was identified, substitution of Lys42 by Arg results in resistance to streptomycin. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bodmer T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              streptomycin; antibiotic; susceptibility; sensitive; resistant; rpsL; mutant; small ribosomal subunit; S12 ribosomal protein; polymerase chain reaction; Mycobacterium tuberculosis; ss.
DNA. The spheroidin gene is naturally expressed at high levels. Small
                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rapid detection of antibiotic resistance in Mycobacteria - esp. isoniazid, rifampicin or streptomycin resistance in tuberculosis by detecting mutation in katG, rpoB or rpsL genes.
                                                                                                                                                                                                                ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer ML51 to amplify M.tuberculosis rpsL gene fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Telenti A,
                                                                                                                                                              0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                  Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Honore N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zhang Y,
                                                                                                                                                                                                                                                               3715 GAGGIGICACCCAAACCGGC 3734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (INSP ) INST PASTEUR.

(WEDI-) MEDICAL RES COUNCIL.

(ASSI-) ASSISTANCE PUBLIQUE.

(UYPB-) UNIV CURIE PARIS VI P & M.

(UYBB-) UNIV BERNE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 51; 97pp; English
                                                                                                                                                                                                                                                                                          20 GAGGTGTTACCCAACCAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-00929206.
92FR-00011098.
93FR-00004545.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cole S, Young D,
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ61458 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                       Ouery Match
Best Local Similarity 85.0°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1993-368812/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-AUG-1992;
17-SEP-1992;
16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-APR-1993;
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17-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ61458;
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                                                                                                                                                                                                                                                                                                                                                                               AAQ61458
AXC AAQ61458
AXC AAQ61458
AXC AAQ61458
DT 25-MAR-2
DT 17-MAY-1
DX BLTEPLON
WW STREPCON
WW WO932245
WW WOSSELLIA
WW WDI, 198
WW WD
888888
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                                                                                                                                                                                                                                                                                                                                New protein tyrosine kinase and related nucleic acid - vectors, transformed cells, etc., useful for diagnosis and treatment of tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Three primers (AAQ64159-Q64161) were used to amplify regions of the protein tyrosine kinase receptor. The gene encoding the receptor is related to the trk proto-oncogene. Antibodies against the encoded polypeptide are useful for diagnosis and for the treament of tumours. The antibodies may also be radiclabelled or coupled to a cytotoxin for destruction of cancer cells. Antisense nucleic acid can be used to inhibit gene expression. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                      Holtrich U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                         (CHEM-) CHEMOTHERAPEUTISCHES FORSCHUNG
                                                                                                                                                                                                                                      Strebhardt K, Ruebsamen-Waigmann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1798 AGTGACGTCTGGTCCTTTGG 1817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 AGTGATGTCTGGACCTATGG 1
                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 8; 9pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
                                                                                            92DE-04239817.
                                                                                                                                         92DE-04239817
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis B primer MD 123.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17; Conservative
                                                                                                                                                                                                                                                                                      WPI; 1994-184380/23
DE4239817-A1
                                                                                            26-NOV-1992;
                                                                                                                                            26-NOV-1992;
                                             01-JUN-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ45172;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1282
AAQ45172/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Loca
Matches
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0.4%; Score 15.2; DB 1; Length 20;

Gaps

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New entomopoxvirus polynucleotide sequences, proteins and vectors - are used for expression of heterologous proteins in both insect and mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA from human embryonic tissue was isolated. With the use of primer P6(4) PTK-specific cDNA was synthesised. The cDNA was amplified using primers P6(4) and NS. A 2097 bp DNA fragment was obtained. Primers E3, P12 and E6 were then used in the isolation of the C-terminal of the HEKZ receptor gene. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human embryonal kinase, HEK; protein tyrosine kinase, PTK; tumour;
cancer; therapy; amplification; primer; polymerase chain reaction; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human embryonal kinase 2-receptor protein - useful in tumour diagnosis
                                                                                                                                                  An AmEPV clone spanning a central region of the spheroidin gene was isolated by PCR amplification of AmEPV genomic DNA using the primers given in AAQ66802-03. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ä
                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                     0.4%; Score 15.2; DB 1;
85.0%; Pred. No. 1.5e+03;
ative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Holtrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHEM-) CHEMOTHERAPEUTISCHES FORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ruebsamen-Waigmann H,
                                                                                                                      Disclosure; Page 75; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1798 AGTGACGTCTGGTCTTTGG 1817
                                                                                                                                                                                                                                                                                                                         3715 GAGGTGTCACCCAAACCGGC 3734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 10; 11pp; German.
                                                                                                                                                                                                                                                                                                                                                20 GAGGTGTTACCCAACCAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                2462/c
AAQ62462 standard; cDNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92DE-04233782.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HEK-2 receptor primer P6(4)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                   Query Match.
Best Local Similarity 85.04
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 85.0°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Strebhardt K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-OCT-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-NOV-1994
                                                                   used for exhost cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ62462;
                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1284
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Amplification of nucleic acid from a blood sample by enzymatic amplification is new. No prepn. of the blood sample otherwise necessary to prepurify the nucleic acid sequences to be amplified is performed and the proportion of the sample in the reaction mixt. is greater than 5 vol. if a specific amt. of salt is present in the reaction mixt. The new process was exemplified using HLA DQ, Factor IX, Hepatitis B and Rubella DNA and primers: (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                opt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                           Amplification and detection of nucleic acid in unprocessed blood - contg. anticoagulant, in presence of controlled concn. of salt, for detecting genetic sequences or microorganisms.
HLA DQ; Factor IX; Hepatitis B; Rubella; detection; amplification;
primer; polymerase chain reaction; PCR; blood; anticoagulant; salt;
enzymatic amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Spheroidin; gene expression; vector; insect cell culture; mammal cell culture; AmBPV; Amsacta moorei; entomopoxvirus; PCR; polymerase chain reaction; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 0.4%; Score 15.2; DB 1; Length 20; Local Similarity 85.0%; Pred. No. 1.56+03; Local 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 9 A; 2 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                          (HOFF ) HOFFMANN LA ROCHE & CO AG F.
                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure, Page 9, 29pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         925 TICCIGITCAICCIGGIGGT 944
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ66802 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-00991867
                                                                                                                                                                           93EP-00113861
                                                                                                                                                                                                            92CH-00002875
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYFL ) UNIV FLORIDA
                                                                                                                                                                                                                                                                                                             WPI; 1994-110893/14.
                                                                                                                                                                           31-AUG-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-DEC-1992;
                                                                                                                                                                                                            11-SEP-1992;
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18-JAN-1995
                                                                                                        EP590327-A2
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                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ66802;
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Gaps

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ricinoleic acid;

vivlemore401-10.rng

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Prodn. of hydroxylated fatty acids, e.g ricinoleic or lesquerolic acid in genetically modified plants such as rapeseed, flax, sunflower or cotton, contg. a fatty acid hydroxylase gene.
                                                                            tränsgenic plant; oilseed; seed oil; rapeseed; Arabidopsis; Crambe; Brassica juncea; canola; flax; sunflower; safflower; cotton; cuphea; soybean; peanut; coconut; oil palm; corn; fahl2; castor; Ricinus communis; primer; PCR; polymerase chain reaction; ss.
                     Kappa fatty acid hydroxylase fah12 gene PCR primer HR1.
                                                          Kappa fatty acid hydroxylase; lesquerolic acid;
                                                                                                                                                                                                                                                                                        95WO-US011855.
                                                                                                                                                                                                                                                                                                                            94US-00314596.
                                                                                                                                                                                                                                                                                                                                             94US-00320982.
95US-00530862.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Somerville C, Broun P,
                                                                                                                                                                                                                                                                                                                                                                                                                                        (VLOO/) VAN DE LOO F J.
                                                                                                                                                                                                                                                                                                                                                                                                    (SOME/) SOMERVILLE C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-200914/20.
                                                                                                                                                                                                                                                                                                                                                                                                                          BROUN
                                                                                                                                                                                                              WO9610075-A1
                                                                                                                                                                                                                                                                                        25-SEP-1995;
                                                                                                                                                                                                                                                                                                                            26-SEP-1994;
                                                                                                                                                                                                                                                                                                                                               .1-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                 20-SEP-1995;
                                                                                                                                                                                                                                                   04-APR-1996,
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14-MAY-1997
                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1287
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This antisense primer is derived from exon-5 of the human retinal rod CGMP-phosphodiesterase beta-subunit (PDE-beta) gene, and may be used for PCR amplification and sequencing of normal and mutant forms of the PDE-beta gene. The primer may be used along with sense primer AAT17129 or AAT17135 to detect a variant with a C-to-T transition at position 11638 are xon-5, resulting in a nonsense mutation (GHD298X) (e.g. in patients AR77 and AAI20) linked with autosomal recessive retinitis pigmentosa. Mutations in the rhodopsin and retinal degradation slow protein genes are also implicated in retinitis pigmentosa. Detection of any of these mutations in a foetus or patient may be used in diagnosis. (Updated on 25-MAR-2003 to correct PP field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diagnosis of hereditary retinal degenerative diseases e.g. retinitis pigmentosa, - caused by a human photoreceptor protein mutation, by detection of the mutation by PCR amplification or hybridisation methods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                      Primer; human; cGMP-phosphodiesterase; beta-subunit; mutation; PCR;
                                                                                                                                                                                                                                                 Primer for cGMP-phosphodiesterase beta-subunit gene amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .,
                                                                                                                                                                                                                                                                                                          polymerase chain reaction; eye; rod; retina; photoreceptor; retinitis pigmentosa; diagnostic; prenatal diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 12 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              319 CCCACTCCCTCCATCTCCTG 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 9; Col 15; 71pp; English
                       AGTGATGTCTGGACCTATGG 1
                                                                                                               AAT17130 standard; DNA; 20 BP
93US-00033081
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90US-00469215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91US-00805123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT27015 standard; cDNA; 20
                                                                                                                                                                                          (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HARD ) HARVARD COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dryja TP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-159684/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-DEC-1991;
                                                                                                                                                                                          25-MAR-2003
03-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                    US5498521-A
                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Berson EL,
                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                     20
                                                                                                                                                     AAT17130;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                              RESULT 1285
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1D AAT270:
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AC AAT270:
XX
DT 25-MAR-
DT 10-SEP-
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Van De Loo FJ;

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                                        PCR primers HF2 (AAT27014) and HR1 (AAT27015) were designed to amplify a 700 bp fragment of the kappa fatty acid hydroxylase fally gene of castor. PCR amplification was used to detect the fall2 sequence in Arabidopsis plants transformed, by Agrobacterium tumefaciens-mediated transformation, with the fall2 gene on binary Ti plasmid pBG. Transgenic plants were identified that showed altered levels of hydroxylated fatty acids, e.g.
                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein kinase; treatment; disorder; cancer; human; primer; PCR; foetal liver; degenerate; polymerase chain reaction; ss.
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0
                                                                                                                                                                                                                                                                                0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Degenerate PCR primer for protein kinase cDNA isolation.
                                                                                                                                                                                                                                         Seguence 20 BP; 6 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
Example 1; Page 38; 105pp; English
                                                                                                                                                                                                                                                                                                                                                                         2982 CAGGGCTTTTTCTGGCACCG 3001
                                                                                                                                                                                                                                                                                                                                                                                                                 20 caagecerrircreeracce 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT38287 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first entry)
                                                                                                                                                                                                                                                                                                       Local Similarity 85.0
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(revised)
(first entry)

25-MAR-2003 10-SEP-1996

AAT27015;

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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           Ricinus communis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9730582-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Broun P,
                                                                                                                                                                                                                                                                                                            AAT95010;
                                                                                                                                                      Query Match
*888888888888
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                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                              The present sequence is a degenerate PCR primer for protein kinase (PK) cDNA isolation. The PK may be used to treat disorders related to abnormal PK activity, and cancer. Human foetal liver polyA+ RNA was PCR amplified, and the products subcloned, sequenced and used to produce the plasmids pLKB1-1 and pLKB1-2. When the coding region of pLKB1-1 was isolated by restriction digest, and analysed by northern blotting against polyA+ RNA prepared from various human organs and cultured cells, weak expression was detected in almost all tissue and cell types. (Updated on 25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New oligonucleotide primers amplifying gene regions conserved among mammals - useful for developing genomic maps, isolating clones and making cross-species comparisons.
                                                                                                                                        DNA encoding protein kinase - for potential treatment of protein kinase activity related disorders and cancer.
                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer; polymerase chain reaction; amplification; UM-STS; universal mammalian sequence tagged site; genomic map; clone;
                                                                                                                                                                                                                                                                                                                                             0.4%; Score 15.2; DB 1; Length 20; 70.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                     4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homeobox 7 PCR primer for universal mammalian STS's.
                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 1 C; 4 G; 4 T; 0 U; 6 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yuzbasiyan-Gurkan V;
                                                                                                                                                                                                                                                                                                                                                                     2; Mismatches
                                                                                                                                                                            Example A-1; Page 23; 30pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                            1288 GTAGCCGTGAAGATGCTGAA 1307
                                                                                                                                                                                                                                                                                                                                                                                                                  1 GTNGCNGTNAARATGYTNAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 9; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV01150 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97WO-US002403.
                      96WO-JP000660
                                             95JP-00057104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                   (CHUS ) CHUGAI SEIYAKU KK
                                                                                                                                                                                                                                                                                                                                                                   14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brewer GJ, Venta PJ,
                                                                                                                                                                                                                                                                                                correct PA field.)
                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-435083/40.
                                                                                                                 WPI; 1996-433826/43.
                      15-MAR-1996;
                                             16-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV01150
                                                                                            Nezu J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1288
                                                                                                                                                                                                                                                                                                                                                                     Matches
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8 g

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The present sequence represents a specifically claimed oligonucleotide PCR primer. The oligonucleotide can be used for polymerase chain reaction (PCR) amplification of DNA, specifically regions of specific genes that are conserved among mammalian species, i.e. pairs of oligonucleotides from the present specification represent universal mammalian sequence-tagged site (UM-STS) primers. The primers are used to develop genomic maps, to isolate clones from libraries, to make cross-species comparisons and to develop additional genetic markers. UM-STS allow genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primers HR1 (AAT95010) and HF2 (AAT95009) were designed to amplify a 700 bp fragment of the castor kappa hydroxylase fahl2 gene. They were used to verify the presence of an intact fahl2 gene in young leaves of Arabidopsis thaliana plants that had been transformed, via Agrobacterium-mediated transformation, with the castor fahl2 gene on binary Ti plasmid plass. Transgenic Arabidopsis plants produced richnoleic acid, leaquerolic acid, densipolic acid and auricolic acid. The invention relates to the production of hydroxylated fatty acids by genetically modified plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kappa hydroxylase, fatty acid hydroxylase, fah12 gene,
fatty acid unsaturation, transgenic plant, vegetable, seed oil, oilseed,
castor, primer, PCR, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Altering the amount of unsaturated fatty acid in plant seeds - by g manipulation of fatty acid desaturase or hydroxylase, particularly increase oleic acid content.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ပဲ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Somerville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Castor kappa hydroxylase fah12 gene PCR primer HR1.
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 8 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Boddupalli SS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 39; 119pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1348 GAGATGGAGATGATGAAGAT 1367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CARN-) CARNEGIE INST WASHINGTON (MONS ) MONSANTO CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 GAGCTGGAGAAGCTGAAGAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT95010 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 85.0
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-434749/40.
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vivlemore401-10.rng

BP.

AAV85579 standard; DNA; 20

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Gaps

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10-FEB-1999 (first entry) LRP5 PCR primer GI 1F. LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzhe, disease; low density lipoprotein;

PCR primer; ss

Homo sapiens. WO9846743-A1

Synthetic.

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This antisense oligonucleotide is specifically hybridisable with selected DNA or RNA deriving from the puromycin-sensitive aminopeptidase (PSA) -99. This oligonucleotide is used for dignosing conditions associated with PSA expression. The human PSA-99 (875 amino acids) and the murine PSA-99 (920 amino acids) both exhibit PSA activity and can be used to generate anti-PSA antibodies. Cell lines which produce the antibody and host cells transfected with vector containing nucleic acid molecules encoding the PSA and the oligonucleotides can be used in assays for identification of agents which act by targeting PSA, for modulating PSA activity or endogenous PSA substrates, proliferation rate or viability of cells or to induce apoptosis within cells by inhibiting PSA activity. Agents which can diminish PSA activity in cells, by modulation of the amount of PSA in cells due to modulation of PSA ymphesis, may be used in treatment of proliferative diseases, including tumours such as leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated protein with puromycin-sensitive aminopeptidase activity - which may be used in treatment of proliferative disorders, including cancer and
                                                                                                                                                                                                                                                                                                          Puromycin-sensitive aminopeptidase; PSA; traatment; cancer; psoriasis; proliferative disorder; hybridise; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                          Puromycin-sensitive aminopeptidase (PSA) antisense oligonucleotide 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tobler AR, Altmann K, Schlapbach R;
 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 3 C; 1 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          carcinomas or epithelial disorders like psoriasis
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 36; Page 116; 141pp; English.
                                    2982 CAGGGCTTTTTCTGGCACCG 3001
                                                       20 CAAGGCGTTTTCTGGTACCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96WO-EP001518.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96WO-EP001518.
                                                                                                                                                              AAV06269 standard; DNA; 20
                                                                                                                                                                                                                                     (first entry)
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fontana A, Constam D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NOVS ) NOVARTIS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-512727/47.
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                         WO9738114-A1
                                                                                                                                                                                                                                       22-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-OCT-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           psoriasis.
                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                   AAV06269;
                                                                                                                            RESULT 1290
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Gaps
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                                                                                                                                                                                                                                                                                                   can also be used for detection, diagnosis and drug screening
                                                                                                                                                                                                                                                                                                                        Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                               157 GCTCCATCCTCGGGAGATGA 176
                                                                                                                                                                                                                                                                                                                                                          20 GCTGCATCCTCTGGAGAGA 1
                                                                                                                                                                                                                                                                                                                                                                                 1510/c
AAV14510 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                AAV14510;
                                                                                                                                                                                                                                                                                                                                                                           RESULT 1292
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ID AAV14
XX
AC AAV14
XX
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Gaps

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ch 0.4%; Score 15.2; DB 1; Length 20; Il Similarity 85.0%; Pred. No. 1.5e+03; 17; Conservative 0; Mismatches 3; Indels

Best Local Similarity

Matches

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Query Match

RESULT 1291 AAV85579/c

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The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV8555 to AAV8556 represent PCR primer used for obtaining LRP5 colba. Nucleic acid molecules (NAMs) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMs or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              presentation, cytokine clearance and/or inflammation, viral infection, pathoganic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention
Claim 12; Page 98; 200pp; English.
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New isolated LDL-receptor related protein - used to develop products for treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.

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Hammond

Hess JW, Caskey CT, Cox RD, Gerhold D, Hammond awaguchi Y, Merriman TR, Metzker ML, Nakagawa Y; 4S, Twells RCJ;

Todd JA, Heb. P. Kawaguchi Y Twel

Hey P, Kawagu Phillips MS,

WPI; 1998-594573/50.

(WELL) WELLCOME TRUST LTD. (MERI) MERCK & CO INC.

98WO-GB001102. 97US-0043553P. 97US-0048740P.

S-APR-1998; 15-APR-1997; 05-JUN-1997;

22-OCT-1998

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This sequence is a probe used to test the method of the invention. The method is for the detection of a hybrid of a target nucleic acid and a probe nucleic acid in a sample comprises: (1) preparation of a labelled cunit composed of late and labelled substances capable of interactive action in a phenomenon of electron flow in double helix structure in the presence of a hybrid via a linker; (2) addition of a probe nucleic acid having a complementary base sequence to the base sequence of target nucleic acid (3) placing the sample under conditions capable of forming the target nucleic acid and the probe nucleic acid to give a hybrid; (4) mixing the sample capable of forming the hybrid and the labelled unit; (5) detection of the change in the lat and/or labelled substances caused by their interaction; and (6) detection of the presence of the hybrid in the sample. The method can be used for the detection of a target nucleic acid having a specified base sequence. The method allows for the simple and easily operable detection of labelled nucleic acids for respective base sequence
                                                                                 Detection of a target nucleic acid - using a labelled unit with labels capable of interactive action in a phenomenon of electron flow in double helix structure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spheroidin, Entomopoxvirus, expression system, replication, heterologous gene expression; thymidine kinase, poxvirus, vaccinia, swinepox virus, insect pest control; immunity, PCR primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer RM92 for Amsacta moorei entomopoxvirus gene sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 11 A; 0 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                χ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2832 ATATATATATATACATATA 2851
                                                                                                                                                                                   Example 3; Page 5; 9pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 ATATATATATATATA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gruidl ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91US-00657584.
92US-00827685.
92WO-US000855.
92US-00991867.
93US-00107755.
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Amsacta moorei entomopoxvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95US-00544332
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ID AAZ10086 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
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Best Local Similarity 85.09
Marches 17; Conservative
                                                           WPI; 1999-350323/30.
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                     (CANO ) CANON KK.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAR-2003
28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-FEB-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-FEB-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US5935777-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ10086;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence is a primer for the Amsacta moorei entomopoxvirus (AmEPV) spheroidin gene. The amplified sequence is an example of the gene of the invention, which encodes a 115 kDa protein. EPV spheroidin and thymidine kinase promoters can be used in DNA constructs and vectors for expression of heterologous genes in insects or mammalian cells, e.g. vectors containing Bacillus thuringiensis toxin genes for use in insect control, or recombinant vaccinia or swinepox viruses for use as viral vaccines. (Updated on 27-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - used in DNA constructs and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                  Entomopoxvirus; spheroidin gene; AmEPV; thymidine kinase; promoter; insect control; viral vaccine; PCR primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vectors for expression of heterologous genes in, e.g. insect cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 0.4%; Score 15.2; DB 1; Length 20; Local Similarity 85.0%; Pred. No. 1.5e+03; les 17; Conservative 0; Mismatches 3; Indels
                                                              Primer RM92 for AmEPV entomopoxvirus spheroidin gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Probe used to test nucleic acid detection method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New Entomopoxvirus nucleic acid sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3715 GAGGTGTCACCCAAACCGGC 3734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 5; Col 22; 55pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid detection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYFL ) UNIV FLORIDA RES FOUND
                                                                                                                                                                Synthetic.
Amsacta moorei entomopoxvirus.
                                                                                                                                                                                                                                                                                                                                                  91US-00657584.
92US-00827685.
92WO-US000855.
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                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-168476/15.
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30-JAN-1992;
12-FEB-1992;
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27-AUG-2003
20-MAY-1998
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New antisense compounds used to treat eg. hyperproliferative conditions.

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                                                                                                                                                                                                                                                                                                                                            virus. The expression vectors can also be used for the control of pests through the insertion of a gene encoding an insect toxin into the expression vector which will infect the target pest and produce large quantities of the toxin. Spheroidin and thymidine kinase are nonessential proteins which makes them ideal for the insertion of exogenous DNA and they are capable of operating in a vertebrate poxvirus (e.g. vaccinia)-mammalian cell-mediated and humoral immunity. (Updated on 20-MAR-2003 to correct PR field.)
                                                                                                                                        PCR primers AA210086-87 were used to amplify Amsacta moorei entomopoxvirus gene sequences. These DNA sequences are used to make expression systems of the invention. The specification describes an Entomopoxvirus (BPV) expression system that is capable of directing the replication and expression of a heterologous gene in a selected host cell. The expression of a heterologous gene in a selected host linked to the selected heterologous gene sequence. The expression system is used for the expression of heterologous sequences and the production of selected proteins in insect and mammalian host cells e.g. human, rodent and primate cells. BPV thymidine kinase and spheroidin genes can also be used in vertebrate poxviruses such as vaccinia and swinepox
                                                   expression system for the expression of heterologous sequences in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mdm2 gene; proliferation; tumour; phosphorothioate; p53; cancer; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; lung cancer; soft tissue cancer; psoriasis; fibrosis; atheroselerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mdm2 phosphorothioate oligodeoxynucleotide #147.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.2; DB 1;
85.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3715 GAGGTGTCACCCAAACCGGC 3734
                                                                                                       Example 5; Col 26; 72pp; English
                                                                       insect and mammalian host cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 GAGGTGTTACCCAACCAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ37617 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17; Conservative
                  WPI; 1999-457596/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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Cowsert LM;

Graham MJ, Monia BP,

Nero P,

Miraglia LJ,

WPI; 1999-610754/52

(ISIS-) ISIS PHARM INC

99WO-US006702 98US-00048810

26-MAR-1999; 26-MAR-1998;

30-SEP-1999,

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                                                                                     AAZ3741, AAZ3742, AAZ37739, AAZ37741 and AAZ37741 the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of hyperproliferative gene expression. The antisense compound is used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compounds used to treat eg. hyperproliferative conditions.
                                                                         AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mdm2 gene, proliferation, tumour; phosphorothioate; p53; cancer; antisense; modulation; oligonuclactide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; hyperproliferation; blood cancer; psoriasis; fibrosis; atherosclerosis; lung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cowsert LM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mdm2 phosphorothioate oligodeoxynucleotide #209.
                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monia BP,
                                     Example 9; Page 51; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                          1346 CTGAGATGGAGATGATGAAG 1365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 9; Page 53; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 CTCAGATGAAGATGATGAGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ37679 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US006702.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-00048810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 85.0'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Miraglia LJ, Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-610754/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9949065-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ37679;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1296
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vivlemore401-10.rng

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Detection, metastasis, melanoma, breast cancer, C-Met, GalNAc, MAGE-3;
CK20, beta-HCG, MAGE-1, melanoma marker gene, tyrosinase, MART-1, MUC-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting melanoma and breast cancer, detecting subclinical metastasis and monitoring treatment involves detecting the presence or absence of nucleic acid targets in the obtained patient's biological sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NGIC-) NGI CANCER TECH CO LLC.
                                                                            AAA39238 standard; DNA; 20 BP.
                                                                                                                                                                                                                               C-Met PCR primer SEQ ID NO:27.
                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                   metastatic; tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hoon DSB, Schmid P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-349569/30.
                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-MAR-1995;
                                                                                                                                                                           07-SEP-2000
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                                                                                                                           AAA39238;
                            RESULT 1298
                                                    AAA39238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a PCR primer used to isolate DNA encoding the human fatty acid desaturase, FADS-2, of the invention. An antibody directed against the 3 FADS molecule of the invention (FADS-1, FADS-2, and FADS-3) is useful for diagnostic or therapeutic purposes. The FADS coding sequences are useful in gene therapy. The polypeptide and antibodies are useful in screening for modulating drugs. The polypeptides are also useful for treating liver disease, coronary artery disease and cancer. Note: Two copies of the sequence listing are present within this patent, which contain different sequences. AAA90952 and AAA80955 are both as being SEQ ID's 7-22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel cDNA molecules encoding three human fatty acid desaturases, FADS1, FADS2 and FADS3, useful in the treatment of liver disease, coronary artery disease and cancer.
treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, fatty acid desaturase, FADS-1, FADS-2; FADS-3; gene therapy; liver disease, coronary artery disease, cancer, PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human fatty acid desaturase 2 coding sequence PCR primer TU13-F1.
                                                                                                                                                                                                                               ö
                                                                                                                                                                           Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
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                                                                                                                           Sequence 20 BP; 11 A; 0 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 8 A; 8 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                               3462 TTATATATATCTATATAT 3481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 13; Page 7; 72pp; English.
                                                                                                                                                                                                                                                                                                                                 20 rrarararrrcraacrarar 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MULT-) MULTIGENE BIOTECH GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA90967 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99EP-00104664.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA90967;
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                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1297
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XA AAA9
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97US-00987326. 95US-00406307.

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mentoring treatment in a patient by amplifying nucleic acid targets monitoring treatment in a patient by amplifying nucleic acid targets (NT) if present in obtained patient's biological sample, and detecting the presence of NT which is indicative of the presence of metastatic melanoma and breast cancer cells. Also described is a kit useful for detecting melanoma and breast cancer cells comprising a pair of primers for amplifying NT from a set of breast cancer marker genes which includes C Met, GalNAc, MAGE-3, CKZO, betaHCG and MAGE-1 or a set of melanoma marker genes including tyrosinase, MART-1 and MAGE-3 and containers for each pair of primers. The method is useful for detecting metastatic melanoma cells and breast cancer cells, subclinical metastasis and for monitoring treatment in the patient's biological sample such as blood, tumour draining lymph node, bone marrow and cerebrospinal fluid. The method is sensitive, specific and detects occult melanoma or breast cancer cells in the blood of patients with or without clinical evidence of the disease.

The present sequence represents a PCR primer for C-Met, which is used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                     The present invention describes a method for detecting (D) metastatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 20 BP; 0 A; 7 C; 5 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1809 GTCCTTTGGGGTCCTGCTCT 1828
Example 17; Col 42; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 crecriridecercercrer 20
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Gaps

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3; Indels

0; Mismatches

Conservative

17;

Best Loca Matches

2458 GAGGGCCTTTGTTCTGGGG 2477

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20 GATGGGCTTTTGTTCTGAGG 1

26-JUL-2002 (first entry)

Nucleic acid probe m.

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MEK kinase 5; MAP/ERK kinase kinase 5; ASK1; pro-apoptotic; apoptosis signal-regulating kinase 1; programmed cell death; serine/threonine kinase; MAP kinase cascade; UK/SAPK module, Jun N-terminal kinase/stress-activated protein kinase; p38 module; MKK3; SEK1; transcription factor modulation; expression inhibition; antisense; inflammation; wound healing disorder; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                            Antisense compounds useful for treating or preventing infection, inflammation or tumor formation by inhibiting expression of human MEKK5.
                      Human MEKKS phosphorothioate antisense oligonucleotide, SEQ ID NO:27.
                                           Human MEXKS; mitogen-activated protein kinase kinase b;
                                                                                                                                                                                                                                                                                                                                             Claim 3; Col 39; 35pp; English
                                                                                                                                                                                                 99US-00359757
                                                                                                                                                                                                                       99US-00359757
20-NOV-2000 (first entry)
                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                  Cowsert LM,
                                                                                                                                                                                                                                                                                       WPI; 2000-464034/40.
                                                                                                                                  Homo sapiens
                                                                                                                                                                                                 23-JUL-1999;
                                                                                                                                                                                                                       23-JUL-1999;
                                                                                                                                                       JS6080546-A.
                                                                                                                                                                           27-JUN-2000.
                                                                                                                                                                                                                                                                  Monia BP,
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Gaarde W;

configuration and the mush with the configuration of antibution of antibution. The antisense oligonucleotides were designed to target confiderent regions of the human MEKKS RNA, and were analysed for their effect on MEKKS mRNA levels by district real-time PCR. MEKKS (also known as mitogen-activated protein kinase kinase 5, MEK kinase kinase 5, MEK kinase kinase 6, apoptosis signal-regulating kinase 1, and ASKI) is a dual-specific serine/threonine kinase which mediates cellular responses to mitogenic stimuli by activating both the JUK/SAPK (Jun Northamson) kinase/stress-activated protein kinase) and p38 modules of MAP kinase cascades. MEKKS is thought to play a critical role in the cregulation of apoptosis (programmed cell death) by interacting with other proteins in this cascade and by phosphorylating downstream targets such as MKKS and SERI. MEKKS also participates in another apoptosis-related signalling cascade involving the modulation of transcription factors. Activation and dimerisation of MEKKS is induced by tumour necrosis factor species. Thioredoxin is able to associate with MEKKS and inhibit mEKKS kinase activity and hence MEKKS-dependent apoptosis. The oligonucleotides confitting activity and hence MEKKS-dependent apoptosis. The oligonucleotides confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and traetment of confitting are useful for diagnosis, prevention and anterest and confitting are useful for diagnosis, prevention and confitting are useful for diagnosis, prevention and confitting are useful for diagnosis, prevention and confitting are useful for diagnosis. ions associated with MEKK5 expression, such as inflammation and Sequences AAA61956-A61995 represent phosphorothioate antisense Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other; wound healing disorders

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Gaps
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0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                  17; Conservative
                   Best Local Similarity
   Query Match
                                   Matches
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BP.

AAC79500 standard; DNA; 20

07-FEB-2001 (first entry)

AAC79500;

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3401 ACGGTTTCCAGGGAGGGCC 3420
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ABL57557 standard; DNA; 20 BP.
                                               ABL57557
RESULT 1300
ABL57557/C
ID ABL57557
XX
AC ABL57557
XX
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The invention relates to the determination of the concentration of a nucleic acid target, using a fluorescently labeled probe which produces reduced fluorescence emission when hybridised to the target nucleic acid. The method comprises measuring the reduction in emission caused by hybridisation. The new method is particularly used to quantify target nucleic acids by a real-time polymerase chain reaction, e.g. for quantifying microbial cells in co-cultures or symbiotic systems, for detecting gene mutations or polymorphisms, and for analysing melting curves of target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and no materials are introduced that inhibit amplification by Tag polymerase (so materials are introduced that inhibit amplification by Tag polymerase (so conventional PCR conditions can be used). The specificity of PCR is kept high (amplification of primer dimers is delayed), and the limit of quantitation is reduced. Complex probes are not needed, and amplification can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation confidenced and graphs so more accurate quantitation is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  possible. The current sequence refresents a nucleic acid probe of the invention that was used for investigating the effects or the numbers of G(s) in each target nucleic acid, and the number of G(s) in its
                                                                                                                                                                                                                                                                                                                                                                                                                Kamagata Y, Kurata S, Yamada K, Yokomaku T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                        Concentration; quantification; mutation detection; polymorphic; polymerase chain reaction; PCR; probe; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 12 A; 3 C; 0 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  corresponding invention nucleic acid probe
                                                                                                                                                                                                                                                                                                                                     (NIBI-) JAPAN BIOINDUSTRY ASSOC.
(AGEN ) AGENCY OF IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3473 TATATATATATITGAG 3492
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                                                                                                                                                                                                                                                          20-APR-2000; 2000EP-00108643.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Kanagawa T,
Furusho K;
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                                                                                                                                                                                                                                                                                                  20-APR-1999;
                                                                                                                                                                              EP1046717-A2
                                                                                                                                      Unidentified
                                                                                                                                                                                                                     25-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                      Kurane R,
Koyama O,
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ID AAC795
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This invention relates to antisense compounds 8-30 nucleobases in length transfer to the 5'-untranslated region, translational start site, translational termination region or 3'-untranslational start site, translational termination region or 3'-untranslated region of a nucleic acid encoding a p38 mitogen activated protein Kinase (MAPK), where the antisense oligonuclectides inhibit the expression of MAPK. Sequences AAC79501 represent human p38alpha MAPK and p38beta MAPK cDNA sequences. AAC79501 represent human p38beta MAPK and p38beta MAPK cDNA AAC79501 and AAC79501 and AAC79501 and AAC79501 and AAC79501 and AAC79501 and AAC7951 included in the invention are a p38alpha cDNA sequence AAC79523 and antisense oligonuclectides AAC79523 - AAC79535 isolated from rat tissue. Murine p38beta MAPK cDNA is represented in AAC79537 and antisense oligonuclectides regreting the sequence are given in AAC79538 - AAC79522. The antisense oligonuclectides have antishematic; antistributaticic; antistributaticic; antistributaticic; antistributaticic; antistributaticic; antistributaticic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunosuppressive; cardiant and antiinflammatory activity. The antisense oligonuclectides are useful for inhibiting the expression of p38 MAPK in cells or tissues. The oligonuclectides are used for treating an animal with diseases such as inflammatory or autoimmune diseases e.g. rheumatoid arthritis, or heart disease. The oligonuclectides are also useful for
                                                     Antisense oligonucleotide; p38 mitogen activated protein kinase; MAPK; antirheumatic; antiarthritic; immunosuppressive; cardiant; heart disease; antiinflammatory; autoimmune disease; rheumatoid arthritis; apoptosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense compound targeted to p38 mitogen activated protein kinase inhibits protein kinase and is useful for diagnosing and treating
                                                                                                                                                                                                                                                                                                                                                                                                                                               Mckay R, Popoff I;
                    Human p38alpha antisense oligonucleotide SEQ ID 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 9 A; 8 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammatory, autoimmune and heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 39; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nero PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 GTTAGTGTGTGTGTGT 1
                                                                                                                                                                                                                                                                                                               04-APR-2000; 2000WO-US008794.
                                                                                                                                                                                                                                                                                                                                                           99US-00286904.
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                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-664982/64.
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                                                                                                                                                                                                                       WO200059919-A1.
                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                           06-APR-1999;
                                                                                                                                                                                                                                                                   .2-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monia BP,
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Claim 14; Col 41-42; 39pp; English.

ö 0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; 3; Indels

AAC92851 standard; DNA; 20 BP RESULT 1302

AAC92851;

(first entry) 27-MAR-2001 Human P13 kinase p55 gamma antisense oligonucleotide, SEQ ID NO:34.

Human phosphatidylinositol 3-kinase p55 gamma regulatory subunit;

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PI3 kinase p55 gamma; hp55-gamma; p55-gamma; PIX3R3; p55PIK; signal transduction; downstream effector; receptor tyrosine kinase; insulin receptor; IR; insulin-like growth factor receptor; IGFR; cell growth; differentiation; apoptosis; developmental regulation; alternative splicing; tumour formation; cancer; inflammation; infection; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                      Novel antisense compound targeted to human pl3 kinase p55 gamma specifically hybridizes with and inhibits the expression of human pl3 kinase p55 gamma, useful for modulating the expression of pl3 kinase p55 gamma in cells.
                                                                                                                                                                                                     Ward DT;
                                                                                                                                        99US-00433694.
                                                                                                                                                             99US-00433694.
                                                                                                                                                                                                      Borchers AH, Cowsert LM,
                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                          WPI; 2001-101697/11.
                                                                           Homo sapiens.
                                                                                                                                         03-NOV-1999;
                                                                                                                                                             03-NOV-1999;
                                                                                               US6165790-A.
                                                                                                                   8-DEC-2000
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such as tumour formation, inflammation and certain infections, and allow expression level modulation of the alternatively spliced forms of PI3 Sequences AAC92827-C92906 represent phosphorothicate antisense cinase p55

Gaps ö 0.4%; Score 15.2; DB 1; Length 20; 35.0%; Pred. No. 1.5e+03; Indels Sequence 20 BP; 6 A; 4 C; 5 G; 5 T; 0 U; 0 Other; 0; Mismatches 3200 AGCTGGAGGATCCCCTCCAA 3219 85.0%; Best Local Similarity 85.0 Matches 17; Conservative Ouery Match ò

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AAH56777 standard; DNA; 20 BP RESULT 1303 AAHS6777
YID AAHS
XX
AC AAHS
XX
DT 06-8

AAH56777;

06-SEP-2001 (first entry)

Human; carbonic anhydrase; CA12; genetic disease; antisense target;

therapeutic; ss

WO200161030-A2. Homo sapiens.

23-AUG-2001.

Human carbonic anhydrase (CA12) protein target DNA #3

(first entry)

15-NOV-2001

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The present invention specifically claims AAH56368 to AAH56832 which are antisense oligonucleotides to nucleotide sequences encoding groE. More generally, antisense compounds (I) comprising antisense oligonucleotides of 5-50 bases targeted to a nucleotide sequence encoding groE. More of 5-50 bases targeted to a nucleotide sequence encoding groE. (heat shock protein (HSP)60) (GL) and groES (HSP10) (GS) gene from a microorganism, where the antisense compound is complementary to GL or GS is claimed. (I) have antibacterial, antiviral and expression of GL or GS, is claimed. (I) have antibacterial, antiviral and for inhibiting expression of GL or GS in cells or tissues in vitro. (I) are also useful for inhibiting the growth of a microorganism, or inhibiting the growth of a microorganism, or inhibiting the growth of a microorganism, or inhibiting the growth of a microorganism or to a cell infected with the microorganism, (I). (I) are also useful for treating a mammalian pathological condition mediated by the microorganisms which involves administering to the microorganisms which involves administering to the microorganisms which involves administering to the correct and administering (I) is such that the growth of microorganism is inhibited. The antisense compounds are utilised for diagnostics, to prophylaxis and as research reagents and kitts, e.g., to prevent or delay microbial infections in humans. They are also useful as molecular weight markers. AAH56362 to AAH56857 and AAH56854 represent PCR primers for grow grows encourage the present invention. AAH56855 and AAH56870 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         groES
of the
                                              Antisense oligonucleotide, groE, groEL, groES, inhibitor, growth, microorganism, Escherichia coli, Streptococcus pneumoniae, diagnosis; Streptococcus pyogenes; Staphylococcus aureus; Pseudomonas aeruginosa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel antisense compounds targeting nucleic acid encoding groEL or gene of microorganism, which hybridize with and inhibit expression genes, useful to inhibit growth of microorganism having the genes.
                                                                                                                     antibacterial; antiviral; antiproliferative; antisense therapy
aureus groE operon antisense oligonucleotide SEQ ID NO:425.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            groß nucleotide sequence given in the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                             (GENE-) GENESENSE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 53; 110pp; English
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                                                                                                                                                                                                                                                                                                                                                                               99US-0166249P.
                                                                                                                                                                                                                                                                                                                                20-NOV-2000; 2000WO-CA001347
                                                                                                                                             microbial infection; ss
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                                                                                                                                                                                          Staphylococcus aureus
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                                                                                                                                                                                                                                      WO200136625-A2
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The invention relates to a method for selecting optimal subsequence

Example 4; Page 23; 87pp; English

Selecting optimal subsequence antisense targets for inhibition of mexpression of target mRNA for the therapeutic treatment of genetic

Ju-Seog L;

Bollon AP, Gray DM, WPI; 2001-529916/58

(GRAY/) GRAY D M. (JUSE/) JU-SEOG L.

14-FEB-2000; 2000US-00504653. 14-FEB-2001; 2001WO-US004732.

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antisense targets. The method involves preparing an antisense oligonucleotide capable of inhibiting mRNA expression of target mRNA sequences; as well as antisense oligonucleotides capable of binding DNA. The antisense and antisense oligonucleotides capable of binding DNA agents for the treatment of genetic disease. The present DNA sequence is human carbonic anhydrase (CAL2) protein target DNA related to the invention. Note: The present sequence is shown as DNA in the specification; however, in vivo, this target sequence would be mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense; mdm2; hyperproliferation; cancer; psoriasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 20 BP; 2 A; 9 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human mdm2 phosphorothioate oligonucleotide #145.
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Gaps

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3; Indels

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3115 TITTAATITITAACTIATIG 3134

17; Conservative

Matches

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rrrrrrrrrcaacrrrrg 20

AAD15577 standard; DNA; 20 BP

RESULT 1304

AAD15577

Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03;

0.4%;

Query Match Best Local Similarity

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The hyperproliferative disorder includes cancer or psoriasis
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                                                                                                                                                                     The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranalated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                              Novel antisense compound 8-30 nucleobases in length targeted to a nuclei acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense; mdm2; hyperproliferation; cancer; psoriasis; ss
                                                Cowsert LM;
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                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mdm2 phosphorothioate oligonucleotide #207
                                                Monia BP,
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                                                                                                                                                                                                                                                                                                                                               1346 CTGAGATGGAGATGATGAAG 1365
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                                                Graham MJ,
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                                                                                                                                              Example 9; Col 29; 77pp; English
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98US-00048810
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Best Local Similarity 85.v.
Thea 17; Conservative
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                        (ISIS-) ISIS PHARM INC
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                                                Miraglia LJ, Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nero P,
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                                                                       WPI; 2001-190948/19
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26-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, antiinflammatory, cytostatic, antisense gene therapy, phosphoenol pyruvate carboxykinase-cytosolic, PEPCK-cytosolic, infection, inflammation, tumour formation, phosphorothioate, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is one of a number of antisense compounds of up to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense compound capable of modulating the expression of phosphoenol pyruvate carboxykinase-cytosolic, useful for preventing or delaying infection, inflammation or tumor formation.
                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PEPCK-cytosolic antisense oligonucleotide ISIS 108107.
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                                                             Length 20
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Sequence 20 BP; 11 A; 0 C; 2 G; 7 T; 0 U; 0 Other;
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Local Similarity 85.0%; Pred. No. 1.5e+03;
hes 17; Conservative 0; Mismatches 3;
                                                          Score 15.2; DB 1;
Pred. No. 1.5e+03;
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85.0%; Pred. No. ...
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                                                    Query Match
Best Local Similarity 85.0°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Butler MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-190979/19
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/note= "OTHER= All phosphorothioate linkages, additionally bases 1-6 and bases 15-20 are 2'-0-methoxyethyl bases, and bases 7-14 are deoxynucleotides"

Cowsert LM;

Monia BP,

Graham MJ,

Miraglia LJ, Nero P,

Human, mdm2, hyperproliferative disorder, cancer, psoriasis, atherosclerosis, tumour, cytostatic, anti psoriatic, anti arteriosclerotic, vasotropic, antisense, phosphorothioate, ss.

Location/Qualifiers

Key modified_base

Homo sapiens

/*tag= a /mod_base= OTHER /note= "OTHER= All

02-JAN-2001; 2001US-00752983.

US2001016575-A1.

23-AUG-2001

98US-00048810. 99US-00280805.

26-MAR-1998; 26-MAR-1999;

MIRAGLIA L J. NERO P. GRAHAM M J. MONIA B P. COWSERT L M.

(MIRA/) (NERO/)

(MONI/) GRAH/)

Human mdm2 antisense oligonucleotide 31773.

21-NOV-2001 (first entry)

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Polynucleotides comprising single nucleotide polymorphisms in the human dopamine receptor D2, useful for detecting mutations associated with, e.g. schizophrenia, Parkinson's and myoclonus dystonia.
                                                                                                   Human; dopamine receptor D2; DRD2; polymorphism; allele specific;
drug target isogene; detection; single nucleotide polymorphism; SNP;
genotype; schizophrenia; Parkinson's disease; myoclonus dystonia; MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 10 A; 9 C; 0 G; 1 T; 0 U; 0 Other;
                                                                               Human DRD2 fragment 7 PCR primer SEQ ID NO:239.
                                                                                                                                                                                                                                                                            Duda A, Nandabalan K,
                                                                                                                                                                                                                                                                                                                                                            Example 1B; Page 40; 135pp; English.
                                                                                                                                                                                                                                                        (GENA-) GENAISSANCE PHARM INC.
                   AAF70496 standard; DNA; 20 BP
                                                                                                                                                                                                                                    99US-0144493P.
                                                                                                                                                                                                               19-JUL-2000; 2000WO-US019644
                                                           (first entry)
                                                                                                                                probe; PCR primer; ss.
                                                                                                                                                                                                                                                                            Denton RR,
                                                                                                                                                                                                                                                                                                WPI; 2001-091967/10
                                                                                                                                                                         WO200105832-A1
                                                                                                                                                                                                                                   19-JUL-1999;
                                                                                                                                                     Homo sapiens
                                                           20-APR-2001
                                                                                                                                                                                            25-JAN-2001.
                                       AAF70496;
                                                                                                                                                                                                                                                                             Chew A,
RESULT 1308
         AAF70496/
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Stephens JC;

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The present invention describes polynucleotides comprising single nucleotide polymorphisms (SNPs) in the human dopamine receptor D2 (DRD2). The polynucleotides may be used in assays to detect and characterise polymorphisms in DRD2 that affect its expression and cativity and are involved in disorders such as schizophrenia, Parkinson's and mycolonus (MD). This information would be useful for studying the biological function of DRD2 as well as in identifying drugs targeting this protein for the treatment of disorders related to its abnormal expression or function. Polymorphisms in the DRD2 gene affect the expression of active and functional polypeptides. Therefore it is a divantageous to detect polymorphisms in the DRD2 gene and how those polymorphisms are combined in different copies of the gene. AAF70261 to AAF70309 represent human DRD2 allele specific oligonucleotide primers which are used in the detection of DRD2 polymorphisms. AAF70405 to AAF70452 represent oligonucleotide primers for the detection of the present DRD2 polymorphisms which are given in the exemplification of the present invention. AAF70453 to AAF70453 trepresent equiven in the present invention of the human DRD2 convention.
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ó
                        Gaps
                        .
0
Length 20;
                        3; Indels
Score 15.2; DB 1;
Pred. No. 1.5e+03;
                        0; Mismatches
                                               2319 GTGTGTGTGTGTGTGTGT 2338
                                                                20 GIGITIGIGIATIGGGTGT 1
0.4%;
                        17; Conservative
Query Match
Best Local Similarity
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)448/c AAS29448 standard; DNA; 20

RESULT 1309

Matches

ઠે g AAS29448

AAS29448/ ID AAS2: XX AC AAS2:

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codon region, state to the 5' untranslated region, translation termination condon region, state of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense objection start the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. The antisense objection of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound calcuse hyperproliferation of human cells. The method, which involves the use of the antisense compound; is also useful for detecting the role of mdm2 expression in various cell functions and physiological processes and useful in both clinical research and physiological processes and useful in both clinical research and chiganostic tools AAS29507 represent the human mdm2 antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               An antisense compound, useful for treating e.g. cancer, comprises nucleobases targeted a region (e.g. translation termination codon region) of a nucleic acid encoding human mdm2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to antisense compounds, 8-30 nucleobases in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Pred. No. 1.5e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 11 A; 0 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oligonucleotides of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3462 TTATATATATCTATATAT 3481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 9; Page 17; 81pp; English.
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Gaps

Human mdm2 antisense oligonucleotide 31442.

(first entry)

21-NOV-2001

AAS29386;

AAS29386 standard; DNA; 20 BP

RESULT 1310 AAS29386/c Location/Qualifiers

Key modified_base Homo sapiens.

๙ 1. .20 /*tag=

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Human, epidermal growth factor receptor; hyperproliferative disease;
Herl; antisense; prophylaxis; psoriasis; phosphorothioate backbone;
                                     ;
0
Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                        Human Her-1 antisense oligonucleotide ISIS #122155.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                        1346 CTGAGATGGAGATGAAG 1365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sennett CF, Wyatt JR, Freier SM;
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                                                                                                           20 CTCAGATGAAGATGATGAGG 1
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16. .20
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/mod_base= n
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/mod_base= r
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/mod_base= 1
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                                                                                                                                                                                                  AAD36546 standard; DNA; 20
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                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                               tumour; cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200226758-A1
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                     09-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                    AAD36546;
                                                                                                                                                             RESULT 1311
                                                                                                                                                                                  AAD36546,
                                                                            ò
                                                                                                         셤
                                                                                                                                                                                                                                    The present invention relates to antisense compounds, 8-30 nucleobases in length targeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and chronic myelogenous leukemia. The antisense compound may be and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful for detecting the role of mdm2 expression in various cell functions and physiological processes and useful in both clinical research and adjagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            An antisense compound, useful for treating e.g. cancer, comprises nucleobases targeted a region (e.g. translation termination codon region) of a nucleic acid encoding human mdm2.
                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER /note= "OTHER= All phosphorothioate linkages, additionally bases 1-6 and bases 15-20 are 2'-0-methoxyethyl bases, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                                                             Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cowsert LM;
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Graham MJ, Monia BP,

Miraglia LJ, Nero P,

COMS/) (NERO/) (GRAH/) (/INOW

WPI; 2001-535565/59.

98US-00048810. 99US-00280805.

26-MAR-1998; 26-MAR-1999; MIRAGLIA L J. NERO P. GRAHAM M J. MONIA B P. COWSERT L M.

MIRA/)

02-JAN-2001; 2001US-00752983

US2001016575-A1

23-AUG-2001

Example 9; Page 16; 81pp; English

Sequence 20 BP; 4 A; 7 C; 2 G; 7 T; 0 U; 0 Other;

oligonucleotides of the present invention

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US6399378-B1.
                            04-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                           ABK68247;
                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1313
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                                                                       The invention relates to an antisense oligonucleotide targetted to a nucleic acid molecule encoding human epidermal growth factor receptor (Herl) to inhibit its expression. The antisense compounds are useful for treating diseases or conditions associated with Her-1 such as hyperproliferative diseases especially cancer (lung, ovarian, colon or prostrate cancer) and psoriasis. They are also useful as research reagents, diagnostics, therapeutics, kits and prophylactically e.g. to prevent or delay tumour formation. The present sequence is an antisense oligonucleotide targetted to human Her-1
                 Novel antisense oligonucleotide that specifically hybridizes with and inhibits nucleic acid encoding epidermal growth factor receptor, useful for treating hyperproliferative disease such as cancer or psoriasis.
                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                   Antisense, RECQL2, Bloom's disorder, prophylaxis, infection, tumour; inflammation, therapy, human, phosphorothioate, ss.
                                                                                                                                                                                                          ö
                                                                                                                                                                                     0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                         3; Indels
                                                                                                                                                                                                                                                                                                                                                  Human RECQL2 antisense oligonucleotide, ISIS #137549.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "2'-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              note = "Phosphorothioate backbone"
                                                                                                                                                                    Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                           1671 GATCGCAGACTTCGGGCTGG 1690
                                                      Claim 1; Page 45; 169pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          base= OTHER
                                                                                                                                                                                                                                            20 GATCACAGATTTTGGGCTGG 1
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base= m5c
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/mod_base= m5c
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/mod_base= 1
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/mod_base= 1
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                                                                                                                                                                                                        17; Conservative
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16. .20
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*tag=
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WPI; 2002-394234/42.
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                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                              AAD41768;
                                                                                                                                                                                                       Matches
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Antisense compounds targeted to nucleic acids encoding RECQL2 associated with Bloom's disorder, for modulating RECQL2 expression and treating diseases e.g. tumors associated with expression of the RECQL2 in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to antisense compounds targetted to nucleic acid encoding RECQL2 (gene associated with Bloom's disorder) to inhibit the expression of RECQL2. Antisense compounds of the invention are useful for treating diseases associated with expression of RECQL2, in humans. They are useful for diagnostics, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, inflammation or tumour formation. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human RECQL2 DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 7 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse HYPLIP1 locus specific primer aa049675r1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chatterjee A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypercholesterolaemia; chromosome 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1357 ATGATGAAGATGATCGGGAA 1376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 Argargargargacrgggaa 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Col 45; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK68247 standard; DNA; 20 BP
01-MAR-2001; 2001US-00798096.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bodnar JS, Castellani LW, C
Ohmen J, Ross D, Tafuri S,
                                                                  01-MAR-2001; 2001US-00798096.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-SEP-2001; 2001WO-US028181.
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Best Local Similarity 85.0'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (REGC ) UNIV CALIFORNIA.
                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                   WPI; 2002-535979/57
                                                                                                                                                                                                            Ward DT, Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200220847-A2.
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This invention relates to the cDNA and protein sequences of novel proteins HYPLIP1 or FCHL1 and to sequence variations within these genes that have been shown to be associated with lipid disorders.

Coligonucleotide probes that hybridise to the cDNA sequence are useful for Oligonucleotide probes that hybridise to the cDNA sequence are useful for analysing the expression of FCHL1 by detecting the expression of the mRNA transcript in the sample. A host cell transformed with the cDNA of the invention is useful for producing the protein by recombinant means.

Consecution is useful for preventing a lipid disorder associated with expression of FCHL1 such as familial combined hyperlipidaemia, coronary artery disease, atherogenic lipoprotein phenotype,

Typerapobetalipoproteinaemia, hypertriglyceridaemia, familial dyalipidaemic hypertension, syndrome X, obesity, insulin resistance and hypercholesterolaemia. The cDNA sequence is useful in the diagnosis or prognosis of predisposition to lipid disorders and cancers, and also to identify a molecule which enhances or decreases the HYPLIP1 or FCHL1 activity. The present sequence represents an oligonucleotide primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   specific for the mouse HYPLIP1 locus of the invention. The mouse HYPLIP1
                                               Novel HYPLIP1 and FCHL1 genes and their sequence variations associated with lipid disorder and cancer, useful for prognosis, diagnosis and treatment of lipid disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                      Claim 11; Page 75; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    locus is situated on chromosome 3
WPI; 2002-339808/37.
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ö Gaps ö 0.4%; Score 15.2; DB 1; Length 20; 15.0%; Pred. No. 1.5e+03; ve 0; Mismatches 3; Indels 3651 CTTGCTTGCCTGCAGGGCCA 3670 1 criccarcciccaccacca 20 Query Match 0.4%; Best Local Similarity 85.0%; 17; Conservative g

ВЪ. ABN99692 standard; DNA; 20

Human clusterin inhibiting antisense oligonucleotide 26 (first entry) 16-AUG-2002 Homo sapiens. ABN99692;

Human; antisense inhibition; antisense oligonucleotide; clusterin; hypercholesterolaemia; cardiovascular disorder; ss; hyperproliferative disorder; hyperlipidemic disorder; phosphorothioate backbone; 2'-0-methoxyethyl wing. WO200222635-A1

21-MAR-2002

10-SEP-2001; 2001WO-US028235

11-SEP-2000; 2000US-00659791

(ISIS-) ISIS PHARM INC.

Monia BP,

WPI; 2002-404805/43.

Novel antisense compound targeted to nucleic acid molecule encoding clusterin, useful for treating animal having disease associated with clusterin such as hyperlipidemic disorder, cardiovascular disorder.

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The invention comprises antisense oligonucleotides that are capable of inhibiting expression of the human clusterin gene. The antisense oligonucleotides of the invention are useful for inhibiting the expression of clusterin in cells. The antisense oligonucleotides are also useful for treating an animal with a disease or condition associated with clusterin (e.g. hypercholesterolaemia; cardiovascular disorders; hyperproliferative disorders; and hyperlipidemic disorders). The present DNA sequence represents a clusterin antisense oligonucleotide of the invention. NOTE: The present DNA sequence has a phosphorothicate backbone and also contains 2'-O-methoxyethyl wings
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a compound of 8-50 nucleobases in length targeted to a nucleic acid encoding protein phosphatase 1B (FTP1B), where the compound specifically hybridises with and inhibits the expression of FTP1B (e.g. an antisense oligonucleotide). Also included are (1) a compound of 8-50 nucleobases in length which specifically hybridises with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Compound for inhibiting the expression of protein phosphatase 1B (PTP1B) and for treating diabetes, cancer, or obesity, comprises an antisense oligonucleotide targeted to nucleic acid encoding PTP1B.
                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense; protein phosphatase 18; PTP18; ss; probe; human;
type 2 diabetes; obssity; ovarian cancer; chronic myeloid leukaemia;
hyperproliferative disease; antidiabetic; anorectic; cytostatic;
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                                                                                                                                                                                                                                                                        0.4%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 1.5e+03;
Live 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PTP1B antisense oligonucleotide ISIS 142074.
                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                 489 GCAGACGTACACGCTGGACG 508
              Claim 3; Page 83; 125pp; English.
                                                                                                                                                                                                                                                                                                                                                                                  20 GCAGACGCACATGCTGGATG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JUL-2000; 2000US-00629644.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       blood glucose; gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK85323 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                              Local Similarity 85.0
hes 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WYATT J.
FREIER S M.
MONIA B P.
BUTLER M M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COWSERT L M.
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(FREI/)
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The properties of the control of the
     portion of an active site on a nucleic acid encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 targetting human PTPIB
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0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; 3; Indels Pred. No. 1.5e 0; Mismatches Local Similarity 85.0 les 17; Conservative Query Match Matches

Gaps

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AAH77295 standard; DNA; 20 BP AAH77295; RESULT 1316

g 8

RT-PCR primer 1 for amplification of sAC cDNA. (first entry) 21-FEB-2002 AAH77295
XX
XX
AC
AAH7
XXX
DX
XX
CAC

Rat; soluble adenylyl cyclase; rat sAC; AC; signal transduction pathway; cyclic AMP; cAMP; etimulant; inhibitory; cell growth; cell proliferation; cancer; infertility; diabetes; pathological condition; ophthalmological; blood gas monitoring; cytoseratic; antidiabetic; antiinfertility; RT-PCR primer; ss.

Rattus sp.

WO200185753-A1.

15-NOV-2001.

27-OCT-2000; 2000WO-US029872

11-MAY-2000; 2000US-00568407

(CORR) CORNELL RES FOUND INC.

Levin LR; Buck J, WPI; 2002-041584/05.

New isolated nucleic acid molecule encoding a soluble adenylyl cyclase for qualitatively or quantitatively diagnosing conditions arising from soluble adenylyl cyclase activation.

This polynucleotide sequence represents the RT-PCR primer 1 for amplification of soluble adenylyl cyclase (sAC) cDNA. The invention relates to isolated nucleic acid molecules encoding soluble adenylyl

Example 1; Page 53; 108pp; English.

catactule, Auton mediates cellular responses to nutritional conditions and extracellular signals in organisms from prokaryotes to higher eukaryotes.

CAMP has long been known to exert both stimulatory and inhibitory effects on cell growth and proliferation. The invention is used inhibitory effects on cell growth and proliferation. The invention is used for detect an increase or decrease in the level of sAC in a biological sample from a mammal suspected of suffering such a condition is used for diagnosing the likelihood or onset of, or for monitoring the course and severity of a pathological condition (such as cancer, infertility or diabetes)

cassociated with sAC activation, or deficiency of sAC activation. The isolated polynucleotides of the invention are useful for diagnosing a pathological condition, for monitoring blood gases, for qualitatively or quantitatively diagnosing conditions arising from sAC activation, to detect sAC specific cell lines and to detect sAC positive cells in a patient, in particular a human patient. The advantage of this invention is a new method of screening for a modulator of sAC-induced signalling permits rapid evaluation of a cellular response, and short-circuits cells in a continual assays by detecting individual response in a capture of sacration in pathological assays by detecting individual response in a capture of sacration in the consuming biological assays by detecting individual response. ö Antisense; human Integrin beta 4 binding protein; hIbeta4BP; cytostatic; cell proliferation; cancer; gene therapy; phosphorothioate backbone; ss. signals in sAC-induced signal transduction pathway. Signal transduction assays can be performed with very small amount of material. The polynucleotides and proteins of the invention have cytostatic, cyclase (sAC) proteins. Adenylyl cyclase (AC) is the effector molecule one of the most widely used signal transduction pathways. Its product, cyclic AMP (cAMP), is a nearly universally utilised second messenger Gaps ; 0 antidiabetic, ophthalmological and antiinfertility activities 0.4%; Score 15.2; DB 1; Length 20; 5.0%; Pred. No. 1.5e+03; ve 0; Mismatches 3; Indels Human hIbeta4BP antisense oligonucleotide, ISIS #129486. /mod_base= m5c 16. .20 /*tag= c /mod_base= OTHER /note= "2'methoxyethyl nucleotides" 'note= "2'methoxyethyl nucleotides" note = "Phosphorothioate backbone" Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other; Location/Qualifiers 1998 CAAGCAGCTGGTGGAGGACC 2017 l. .20 /*tag= a /mod_base= OTHER 1 cgagcagcriggriggagarcc 20 mod_base= OTHER mod_base= m5c /mod_base= m5c AAD35760 standard; DNA; 20 BP Local Similarity 85.0%; nes 17; Conservative l. .5 /*tag= b 26-JUL-2002 (first entry) で '*tag= f *tag= *tag= modified base modified_base modified base modified_base modified_base modified base Homo sapiens AAD35760; RESULT 1317 AAD35760/ 886666666666666666666666666666666 ò 셤

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The invention relates to an isolated or substantially pure Von Willebrand factor-cleaving protease (VWF-cp) polypeptide. VWF-cp is useful for purifying VWF which involves providing VWF-cp as a ligand, contacting a solution comprising vWF with the polypeptide ligand under conditions where vWF is bound to the ligand and recovering from the ligand purified VWF. VWF-cp is useful for producing anti-vWF cp polypeptide antibodies which involves immunising an animal with VWF-cp and isolating the anti-VWF cp polypeptide antibodies from the animal. VWF-cp is useful for producing a preparation of prophylaxis and therapy of thrombosis and thrombocytic purpura (TTP), thromboembolic disease such as thrombotic thrombocytopaenia or haemolytic-uraemic syndrome. VWF-cp can also be used for processing plasmatic or recombinantly produced VWF. The invention is useful for construction expression systems and generating transgenic animals which express the polypeptide in vivo. The present sequence is human VWF-cp
                                                                                                                                                                                                                                                                       Novel isolated or substantially purified Von Willebrand factor-cleaving protease, useful for producing preparation for therapy of thrombosis and thromboembolic disease such as thrombotic thrombocytic purpura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mouse; HYPLIP1; FCHL1; familial combined hyperlipidaemia; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lusis AJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
                                                                                                                                   Laemmle B, Gerritsen HE, Furlan M, Turecek P, Schwarz Scheiflinger F, Antoine G, Kerschbaumer R, Tagliavacca Zimmermann K, Voelkel D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   De Jong P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 11 C; 1 G; 5 T; 0 U; 0 Other;
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, Wu C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse HYPLIP1 locus PCR primer #224.
                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 34; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   182 ACGGGGAGGACGAGGCTGAG 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, mouse, HYPLIP1, FCHL1, fa
lipid disorder, PCR, primer, ss.
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Ohmen J, Ross D, Tafuri S,
                   22-NOV-2000; 2000US-00721254.
12-APR-2001; 2001US-0083328.
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Best Local Similarity 85.0
Matches 17; Conservative
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                                                                                                                                                                                                                              WPI; 2002-479950/51.
                                                                                        (BAXT ) BAXTER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200220848-A2.
                                                                                                                                                                                   Zimmermann K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to antisense compounds targetted to a nucleic acid encoding human Integrin beta 4 binding protein (hibeta4BB), which specifically hybridises with the nucleic acid and inhibits its expression. The antisense compounds are useful to prevent or treat diseases associated with hibeta4BP expression, particularly conditions involving aberrant or deregulated cell proliferation (e.g. cancer). The hibeta4BP polynucleotide is used in gene therapy. The present sequence is an antisense oligonucleotide targetted to hibeta4BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound targeted to a region of a nucleic acid encoding human Integrin beta 4 binding protein and that inhibits expression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; Von Willebrand factor-cleaving protease; vWF-cp; therapy; enzyme; transgenic animal; immunisation; thromboembolic disease; preeclampsia; thrombocitc thrombocytic purpura; TTP; Henoch-Schonlein purpura; thrombosis; neonatal thrombosytopaenia; haemolytic-uraemic syndrome; transgenic; anticoagulant; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human Von Willebrand factor-cleaving protease cloning PCR primer, 6561.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 2 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid, for treating e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2093 GTGGCCAGGACACCCCCAGC 2112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Col 45-46; 40pp; English.
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                     /*tag= g
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                                                                                                                                                                                                                                                                                                                         17-NOV-2000; 2000US-00716161.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bennett CF, Freier SM;
                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-370579/40.
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modified base
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AAD39345;

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AC AAD39345
AC AAD39345
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Gaps

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The invention relates to an isolated polynucleotide comprising a sequence variation of a mouse HYPLIP1 CDNA or a human FCHL1 (familial combined hyperlipidaemia) gene. The FCHL1 polynucleotide, the FCHL1 polypeptide or antibody immunoreactive to the FCHL1 polypeptide are useful for treating or preventing cancer associated with expression of FCHL1, as well as for treating lipid disorder. The mouse HYPLIP1 CDNA or human FCHL1 gene are also useful for diagnosing or prognosing a predisposition to lipid disorder and cancer. ABK70902-ABK71303 represent mouse HYPLIP1, human FCHL1 coding sequences and PCR primers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Entomopoxyvirus spheroidin gene; EPV; active promoter; thymidine kinase; virus vector system; vaccinia; swine pox virus vector; ds.
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                                    combined hyperlipidemia) for diagnosing, treating
                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                          0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                   3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic oligonucleotide designated RM92 SEQ ID No 15.
                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                    New mouse HYPLIP1 and human FCHL1 (familial genes and their sequence variations, useful preventing lipid disorders and cancers.
                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                             CTTGCTTGCCTGCAGGGCCA 3670
                                                                                          Claim 11; Page 75; 102pp; English
                                                                                                                                                                                                                                                                                                                                              CTTGCATGCCTGCAGGTCGA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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92WO-US000855.
92US-00991867.
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95US-00544332.
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                                                                                                                                                                                                                                                                                                     Conservative
          WPI; 2002-329882/36
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es 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
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12-FEB-1992;
07-DEC-1992;
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17-OCT-1995;
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New Entomopoxyvirus spheroidin gene sequences isolated from infected Amsacta moorei are useful in expression cassettes to regulate expression of heterologous DNA in virus vector systems.

The invention relates to a novel isolated Entomopoxyvirus (EPV) spheroidin gene polynucleotide which is either an active promoter or an active promoter plus a coding sequence, and is able to direct expression

Example 6; Col 27; 88pp; English.

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               sequences of the EPV spheroidin and thymidine kinase genes are useful in expression cassettes to regulate expression of heterologous DNA in virus vector systems, for example in vaccinia and swine pox virus vectors. This polynucleotide sequence represents a nucleic acid relating to the novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleocide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fluorescently-labeled nucleic acid probes for assaying nucleic acids and their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and
of an operably linked heterologous gene in a host cell. The regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     χ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphism detection; mutation detection; disease diagnosis;
                                                                                                                                                                                                     Gaps
                                                                                      isolated Entomopoxyvirus (EPV) spheroidin gene of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kurata S,
                                                                                                                                                             0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
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                                                                                                                            Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LTD.
                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe poly m for assaying nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 13; Page 62; 152pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kanagawa T, Kamagata Y,
                                                                                                                                                                                                                                        3715 GAGGTGTCACCCAAACCGGC 3734
                                                                                                                                                                                                                                                                          20 caccierracccaaccaccc 1
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03-AUG-2000; 2000JP-00236115.
26-SEP-2000; 2000JP-00292483.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        microbial identification; ss
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                                                                                                                                                       Unery Match
Best Local Similarity 85.0°
Matches 17; Conservative
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microbial identification.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kurane R,
                                                                                                                                                                                                                                                                                                                                                                                                         ABL95904;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOE
                                                                                                                                                                                                                                 Mouse C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:197.
                                                                                                                                                                                                                                                         Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EPB2; LAP; TCF5; CRP2; NFIL6; IL6DBP; NF-M; AGF/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness; oxidative stress response; IL-6 signalling mediator; interlexin-6; carbohydrate metabolism; immunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/BBP) beta, useful in vitro for inhibiting expression of human or mouse C/BBP beta in cells/tissues.
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cytosines are 5-methylcytosine"
 Gaps
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cytosines are 5-methylcytosine"
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       location/Qualifiers
                             3473 TATATATATATTGAG 3492
                                                1 TATATATATTTTTTTGG 20
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                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; ss.
                                                                                                                                      ABL94430 standard; DNA; 20 BP
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17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod
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ABL94430/c
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proliferation and functional differentiation. C/BBP beta (also known as C/BPB2, LAP, TCF5, CRP2, NFIL6, IL6DBP, NF-M, AGP/BBP and Apc/BBP)
Cytimarily regulates hormone responsiveness and oxidative stress responses and is a mediator of IL-6 (interleukin-6) signalling. C/BBP beta is thought to be involved in carbohydrate metabolism, immunity, the Thl response, female fertility and gluconeogenic pathways. C/BBP beta is expressed in the liver, lung, spleen, kidney, brain, and testis, with the highest expression found in the lung. It is also expressed at a higher level in malignant ovarian tissue compared with normal ovarian tissue, and its expression in pancreas is upregulated in response to chronically elevated levels of glucose, indicating that it is involved in the impairment of insulin secretion in type II diabetes. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with C/BBP beta expression, such as cancer (particularly ovarian cancer), tumour formation, diabetes (particularly type II diabetes), infection, or inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a method (M1) for designing capture oligonucleotide probes (II) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, K-ras, PCR primer; probe, capture probe, mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
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                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 1 A; 9 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Capture oligonucleptide Zip ID#1085 oligo #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 GGGGGGGGGGCACCGTGC 1
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Best Local Similarity 85.0
Matches 17; Conservative
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         $$33555555555555555$$
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               selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptide of interest, which comprises using a cost effective medium that does not contain serum or any functional (and/or recombinant) full-length protein. The medium comprises water, plant-derived peptone, osmolality regulator, buffer, energy source, amino acids, lipid source or precursor, source of iron, non-ferrous metal ions and one or more vitamins and cofactors. An example of a protein of interest is human erythropoietin (EPO). The method is useful for producing recombinant proteins of interest. The present sequence is an oligonucleotide used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, erythropoietin, cell culture process, vector; PCR; primer; ss;
EPO; recombinant protein production.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Producing a recombinant polypeptide of interest comprises providing a transformed eukaryotic host cell and a serum-free culture medium and culturing the transformed eukaryotic host cell in the culture medium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method of producing a recombinant
Epstein-Barr virus and polio virus, and parasitic infectious agents
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                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.2; DB 1; Length 20;
Pred. No. 1.5e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mueller D, Unterluggauer
                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neomycin resistance gene oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1644 GCTGGTGACCGAGGACAACG 1663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 18; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 gcregegacagagercaace 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bogner F, Kunert R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACC59624 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                          / Match 0.4%;
Local Similarity 85.0%;
hes 17; Conservative C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                          of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BIOC ) BIOCHEMIE GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-493398/46.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC59624;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zeng S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1324
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method of recovering and purifying recombinant human erythropoietin (rhEpo) from a cell culture medium having host cells comprising removing host cells, cellular constituents and debris from the cell culture medium by performing a procedure comprising centrifugation followed by a depth filtration step and centrifugation. The method is useful for recovering and purifying rhEpo from a cell culture medium. The present sequence is an oligonucleotide shown in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      erythropoietin (rhEpo) from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells,
                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Recovering and purifying recombinant human erythropoietin (rhEpo) fro cell culture medium comprising host cells by the removing host cells, cellular constituents or debris and subjecting one or more fractions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, erythropoietin, recombinant protein production, vector, EPO, host cell line, PCR; primer, ss.
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                                                                                             Length 20;
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                                                                                                                                                            Indels
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                             Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                         Score 15.2; DB 1;
Pred. No. 1.5e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neomycin resistance gene oligonucleotide #2.
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                                                                                                                                                                                                                                                                                          1 AAGCTTGTTGGGAAGCCCTG 20
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                                                                                                0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACC59661 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                   Ouery Match
Best Local Similarity 85.0°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BIOC ) BIOCHEMIE GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Palma N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-505187/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          which contain rhEpo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003045996-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACC59661;
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ID ABT3
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AC ABT3
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Porcine reproductive and respiratory syndrome virus.

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                                                                                    Antiarteriosclerotic; cardiant; vasotropic; antinfective; cytostatic; antiinflammatory; inhibitor; antisense gene therapy; atherosclerosis; short heterodimer partner-1; abnormal; lipid; cholesterol metabolism; cardiovascular disease; infection; inflammation; tumour formation; mouse; antisense; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense oligonucleotide targeted to a nucleic acid encoding short heterodimer partner-1, useful for treating diseases involving abnormal lipid or cholesterol metabolism, e.g atherosclerosis or cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HCV; IBDV; IBV; ILTV; PRRSV; immunogen; virucide; vaccine; transgenic;
polymerase; ORF1b; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                      Mouse short heterodimer partner-1 expression oligo SEQ ID No 76.
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85.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRRSV polymerase protein ORF1b specific PCR primer ED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 8 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 16; Page 96; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2396 GCAGAGGTACCCTGGGTGTC 2415
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                                                                                                                                                                                                                                                                                                                  17-JUL-2002; 2002WO-US023245
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ABZ76428 standard; DNA; 20
                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                              Graham MJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-248161/24.
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                                                                                                                                                                                                                                         WO2003012033-A2
                                                                                                                                                                                                      Unidentified.
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                 12-JUN-2003
                                                                                                                                                                                                                                                                              13-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                              Crooke RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention
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Matches
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2XXXXXXXXXXXXX
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The invention relates to a new orally acceptable immunogenic composition comprising unpurified or partially purified recombinant HCV (Hog cholera virus), IBDV (Infectious Bursal Disease Virus), IBN (infectious bronchitis virus), ILTV or PRRSV (porcine reproductive and respiratory syndrome virus) immunogen expressed in a plant, where the immunogen is expressed in the plant at a level so that upon oral administration of the composition to an animal, an immunogenic response is observed. The composition is useful for producing an immunogenic response against HCV, IBDV, IBV, ILTV or PRRSV antigen. Sequences ABZ76423-28 represent primers, based on the ORPID encoding the polymerase protein, used for PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New orally acceptable immunogenic composition comprising unpurified or partially purified recombinant HCV, IBDV, IBV, ILTV or PRRSV immunogen expressed in a plant, useful for producing an immunogenic response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                p38 mitogen-activated protein kinase; p38 MAPK; phosphorothioate; antisense; antiarthritic; antiinflammatory; kinase inhibitor; human; inflammatory disease; rheumatoid arthritis; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human p38-alpha MAPK oligonucleotide ISIS NO 16505.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
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1..20
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 98; 146pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplification and typing of PRRSV
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07-AUG-2002; 2002US-00311106.
                                                                                           38-AUG-2002; 2002WO-US025126
                                                                                                                                                                                                                                    Zeng F, Leung FC;
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ABX78099 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                against the antigen.
                                                                                                                                                                                                                                                                 WPI; 2003-256496/25
                                                                                                                                                                      LAMD/) LAM D M K.
                                                                                                                                                                                     (ZENG/) ZENG F.
(LEUN/) LEUNG F C.
                             WO2003013598-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
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                                                            20-FEB-2003
                                                                                                                                                                                                                                  Lam DMK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX78099;
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0
                                                                                                                                                                      New antisense compound, useful for preparing a composition for diagnosing, treating or preventing inflammatory diseases, e.g. rheumatoid
                                                                                                                                                                                                                                                    nucleobases in length targeted to a nucleic acid molecule encoding p38 mitogen-activated protein kinase (MAPK). The products of the invention have antiarthritic and antiinflammenty activity, can act as act as kinase inhibitors. The antisense compound is useful for preparing a composition for diagnosing, treating or preventing inflammatory diseases, e.g. rheumatoid arthritis or for use in antisense gene therapy. This sequence represents an antisense oligonucleotide used in a method to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Liver stem; progenitor; pancreatic; insulin; glucose; antidiabetic; PC1; ceal therapy; Southern blot; Pdx1; HlxD2; Isl1; ngn1; NlxZ2; Pax4; NeuroD/beta2; Nkx6.1; insulin 1; insulin 11; glucagon; somatostatin; PP; amylase; elastase; GLUT-2; glucokinase; PC2; PC3; carboxypeptidase E;
                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                        This invention describes a novel antisense compound, which is 8-30
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                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 9 A; 8 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rat insulin I gene amplifying sense primer.
                                                                                                                           Mckay R;
                                                                                                                                                                                                                   Example 2; Col 21-22; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                         2321 GTGTGTGTGTGCGTGTGT 2340
                                                                                                                                                                                                                                                                                                                                                                                                                                         (IXIO-) IXION BIOTECHNOLOGY INC
                                                                                                                           Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-AUG-2000; 2000US-00640101
                                                                             99US-00286904
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-OCT-2001; 2001US-0337446P.
                                                                                                                                                                                                                                                                                                                                                                                                    17; Conservative
                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                          Gaarde WA,
                                                                                                                                                WPI; 2003-089122/08
                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                 inhibit p38 MAPK
                                                                             06-APR-1999;
           US6448079-B1
                                 10-SEP-2002
                                                                                                                           Monia BP,
                                                                                                                                                                                            arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1329
                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                       somatostatin, PP, amylase, elastase, GLUT-2, glucokinase, PC1, PC2, PC3 and carboxypeptidase B (CPE)
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Converting liver stem/progenitor cells to a pancreatic functional cell, useful for treating diabetes, comprises transfecting liver cells with pancreatic development gene and/or culturing with pancreatic differentiation factors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibitor; cytostatic; antirheumatic; antiarthritic; antiangiogenic; antianiflammatory; antisense gene therapy; hyperproliferative disorder; cancer; rheumatcid arthritis; angiogenesis; infection; inflammation; tumour formation; phosphorothioate; 2'-O-methoxyethyl; 2'-MOE; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human VEGFR-1 chimeric phosphorothioate oligonucleotide SEQ ID NO:74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vascular endothelial growth factor receptor 1; VEGF receptor; VEGFR;
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                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytidine residues are 5-methylcytidines
                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 5 C; 7 G; 6 T; 0 U; 0 Other;
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1. .20
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                    1609 AAGTGCATCCACAGGGACCT 1628
                                                                             Example 3; Page 17; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 AAGCGCATCCACAGGGCCAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACC86779 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-AUG-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 85.09
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC86779;
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The present invention describes a compound (C) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding vascular endothelial growth actor teceptor.1 (VBGFR-1), where the compound inhibits the expression of VBGFR-1 and specifically hybridises with the nucleic acid encoding C VBGFR-1 or with an 8-nucleobase portion of an active site on the nucleic acid molecule encoding VBGFR-1. Also described: (1) a composition comprising (C) and acariter or diluent; (2) inhibiting the expression of VBGFR-1 in cells or tissues by contacting the cells or tissues with (C) so that the expression of VBGFR-1 is inhibited; and (3) treating an animal sharing a disease or condition associated with VBGFR-1 is inhibited. (C) has antianglogenic, antirheumatic, antiarthritic, administering (C) to the animal so that the expression of VBGFR-1 is inhibited. (C) has antianglogenic, antirheumatic, antiarthritic, cytostatic and antiinflammatory activities, and can be used in antisense expression of VBGFR-1 and for treating diseases or conditions associated with the expression of VBGFR-1 such as hyperproliferative disorders angiogenesis. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, continuous formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. The present sequence represents a human VBGFR-2 chimeric phosphorothioate, antisense oligonucleotide, which is used in an example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inhibitor; cytostatic; antirheumatic; antiarthritic; antiangiogenic; antiinflammatory; antisense gene therapy; hyperproliferative disorder; cancer; rheumatoid arthritis; angiogenesis; infection; inflammation; tumour formation; phosphorothioate; 2'-0-methoxyethyl; 2'-MOE; ss.
                                                                             New antisense oligonucleotide targeted to a nucleic acid encoding vascular endothelial growth factor receptor-1, useful for diagnosing treating cancer, rheumatoid arthritis, or diseases or conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human VEGFR-1 chimeric phosphorothioate oligonucleotide SEQ ID NO:72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor receptor 1; VEGF receptor; VEGFR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.2; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1610 AGTGCATCCACAGGGACCTG 1629
                                                                                                                                                                                 Claim 3; Page 83; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 AGTGCATTCATCGGGACCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACC86777 standard; DNA; 20 BP
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*tag= a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17; Conservative
                                                                                                                                             involving angiogenesis.
  Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                    WPI; 2003-301004/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified base
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Bennett CF,
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Best Local S
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The present invention describes a compound (C) 8-50 mucleobases in length targeted to a nucleic acid molecule encoding vascular endothelial growth factor receptor-1 (VEGER-1), where the compound inhibits the expression of VEGER-1 and specifically hybridises with the nucleic acid encoding VEGER-1 and specifically hybridises with the nucleic acid encoding VEGER-1. Also described: (1) a composition comprising (C) and a carrier or diluent; (2) inhibiting the expression of VEGER-1 in cells or tissues by contacting the cells or tissues with (C) so that the expression of VEGER-1 is inhibited; and (3) treating an animal having a disease or condition associated with VEGER-1 is inhibited. (C) has antianglogenic, antirheumatic, antiarthritic, administering (C) to the animal so that the expression of VEGER-1 is inhibited. (C) has antianglogenic, antirheumatic, antiarthritic, cytostatic and antiinflammatory activities, and can be used in antisense gene therapy. The antisense compounds are useful for modulating the expression of VEGER-1, such as hyperproliferative disorders (C) e.g. cancer), rheumatoid arthritis, or diseases or conditions involving angiogenesis. The antisense compounds are also useful for disquestics, therapeutics, prophylaxis, e.g. to prevent or delay infection, continualmation or tumour formation, as research reagents and kits, and in distinguishing between functions of various members of a biological pathway. The present sequence represents a human VEGER-2 chimeric formatices of phosphorothicate and tissues oligonucleotide, which is used in an example
                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotide targeted to a nucleic acid encoding vascular endothelial growth factor receptor-1, useful for diagnosing or treating cancer, rheumatoid arthritis, or diseases or conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant protein production, vector; host cell line; erythropoietin; EPO; human; selection agent; selectable marker; PCR; primer; ss.
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cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neomycin resistance gene oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 83; 150pp; English.
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                                                                                                                         12-SEP-2002; 2002WO-US029148.
                                                                                                                                                                      13-SEP-2001; 2001US-00953318.
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Matches 17; Conservative
                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                            involving angiogenesis.
                                                                                                                                                                                                                                                     Bennett CF, Watt AT;
                                                                                                                                                                                                                                                                                                  WPI; 2003-301004/29.
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method for determining the relative copy number of a group of target nucleic acid molecules present in a sample. The method comprises: (1) performing a first PCR in a PCR mixture; (2) performing a second PCR in a PCR mixture; (2) performing a second PCR in a PCR mixture; and (3) quantifying the number of copies of the second target nucleic acid product present in the sample containing the target a group of target nucleic acides in a sample; and (2) determining whether a subject is at risk of acquiring Alzheimer's disease. The method is useful for determining the relative copy number of a group of target nucleic acid molecules present in a sample for determining whether a subject is at risk of acquiring Alzheimer's disease. ACC99520 to ACC99730 represent PCR primer used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Producing polypeptide of interest by culturing hybridoma or transformed host cell which comprise a nucleotide sequence encoding polypeptide, in culture medium free from each of a plant-derived or animal-derived
               Determining the relative copy number of a group of target nucleic acid molecules present in a sample by performing a first or second PCR in a PCR mixture and quantifying the number of copies of the second target
                                                                                                                              The present invention describes a multiplex real-time quantitative PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recombinant protein production; mammalian cell culture; human;
                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 10 A; 6 C; 4 G; 0 T; 0 U; 0 Other;
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KATINGER H.
KUNERT R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neomycin resistance gene oligonucleotide #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          1810 TCCTTTGGGGTCCTGCTCTG 1829
                                                                                                Disclosure; Fig 6; 118pp; English
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28-NOV-2001; 2001US-0333868P.
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                                                                nucleic acid product
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified.
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                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method of producing a transformed evkaryotic host cell that expresses a recombinant polypeptide of interest comprising introducing into a eukaryotic host cell first and second polymuclectide vectors that are integrated into the genome of the host cell. Particular polypeptides of interest include human erythropoietin (EPO). The method is useful in producing host cells that express recombinant polypeptides, such as human erythropoietin, the polypeptides. The present sequence is an oligonucleotide used in the
                                                                                                                                                                                                                                          Producing a transformed eukaryotic host cell (e.g. Chinese hamster ovary cell) that expresses a recombinant polypeptide (e.g. erythropoietin) comprises introducing into the host cell a first and a second
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Multiplex real-time quantitative PCR; PCR primer; copy number;
                                                                                                                                                                               Unterluggauer F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.2; DB 1;
Pred. No. 1.5e+03;
0; Mismatches 3;
                                                                                                                                                                               Kunert R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3576 AAGCITGGAGGGAAGCCGTG 3595
                                                                                                                                                                                                                                                                                                                           Example 1; Page 22; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the polypeptides. The present sec
exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AAGCTTGTTGGGAAGCCCTG 20
                                                                                                                                                                            Schoergendorfer K, Windisch J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACC99708 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%;
                                                                               26-NOV-2002; 2002WO-EP013297
                                                                                                             28-NOV-2001; 2001US-0333868P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-DEC-2002; 2002WO-US038806
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-NOV-2001; 2001US-0336095P
19-JUL-2002; 2002US-0397475P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer SEQ ID NO:89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alzheimer's disease; ss
                                                                                                                                          (BIOC ) BIOCHEMIE GMBH
                                                                                                                                                                                                                                                                                           polynucleotide vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             THERIANOS S.
                                                                                                                                                                                                            WPI; 2003-505205/47
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                WO2003046187-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003048377-A2.
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                                               05-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACC99708;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1333
ACC99708/c
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Gaps

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Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;

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interest, which involves culturing a hybridoma or transformed host cell interest, which involves culturing a hybridoma or transformed host cell in a culture medium for mammalian cell culture, where the medium comprises water, buffer, energy source, animo acids lipid source, or precursor, or iron source, animo acids lipid source, or iron source, animal acids lipid source, animal-derived peptone. An example protein which may be produced using the method is human erythropoietin (EPO). The method is useful for producing a polypeptide of interest chosen from human growth hormone, human monoclonal antibodies, preferably of subclasses 1gG, 1gM and 1gA, and monoclonal human/mouse chimeric antibodies, where the transformed host cell comprises at lass two polymucleotide vectors that encode different proteins of interest and where the different proteins are different parts of an antibody, particularly the light and heavy chains of an antibody. The present sequence is an oligonucleotide used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New primordial human neural crest stem cell having a pluripotent and self -renewing properties, useful for implantation in vivo for cell therapy treatment of human neurological disorders and diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, neural crest stem cell line, transplantation, cell therapy; neurological disease; HNC10 cell, neuroprotrotective; cerebroprotective;
                invention relates to a method of producing a polypeptide of 
nich involves culturing a hybridoma or transformed host cell
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                3; Indels
                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human HNC10 cell TrkB gene PCR primer #4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    3576 AAGCTTGGAGGGAAGCCGTG 3595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 AAGCTTGTTGGGAAGCCCTG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACF04057 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-APR-2001; 2001WO-US013354.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYPE-) UNIV PENNSYLVANIA.
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                                                                                                                                                                                                                                                                                                                                                                                                17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-559151/52.
                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                              Query Match
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The present invention relates to a primordial human neural crest stem cell line suitable for on-demand implantation in vivo into a living host subject comprising a pluripotent and self-renewing neural crest stem cell of human origin. The cell line is useful in the cell therapy treatment of human neurological disorders and diseases. The present sequence is a PCR primer used to isolate human genes from the HNC10 cell line

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  combined hyperlipidemia 1 gene having a variation that is associated with a lipid disorder, useful for identifying susceptibility to the lipid
                                                                                                                                                                                                                                                                                         Mouse; PCR; primer; ss; HYPLIP1; FCHLL; variation; lipid disorder; allele; anti-lipid disorder; anti-cancer therapy; gene therapy; familial combined hyperlipidaemia; coronary artery disease; atherogenic lipoprotein phonotype; hyperapobetalipoproteinaemia; hypertriglyceridaemia; low density lipoprotein subclass B; LDL; familial dyslipidemic hypertension; syndrome X; hypercholesterolaemia; obesity; insulin resistance; cancer; cytostatic; antilipaemic;
                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated polynucleotide comprising a mouse or human familial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lusis AJ;
   Length 20;
                                 Indels
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Score 15.2; DB 1;
Pred. No. 1.5e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chatterjee A,
, Wu C;
                                                                                                                                                                                                                                                              Mouse HYPLIP1 locus PCR primer #230.
                                                               1678 GACTICGGGCTGGCCCGGGA 1697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 11; Page 39; 63pp; English.
                                                                                          20 GACTTTGGGATGTCCCGGGA 1
                                                                                                                                                                      BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-SEP-2001; 2001US-00949428.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUN-2000; 2000US-0213322P
   Query Match 0.4%;
Best Local Similarity 85.0%;
Matches 17; Conservative
                                                                                                                                                                      ADA15290 standard; DNA; 20
                                                                                                                                                                                                                                06-NOV-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CASTELLANI L W. CHATTERJEE A.
                                                                                                                                                                                                                                                                                                                                                                                                     hypotensive, anorectic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LUSIS A J.
OHMEN J.
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TAFURI S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JONG P D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003064372-A1
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                                                                                                                                                                                                    ADA15290;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JONG/
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                                                                                                                                         RESULT 1336
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                                                                                                                                                         ADA15290
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caids span the variation position, a kit for the detection of the FCHLI caids span the variation position, a kit for the detection of the FCHLI course comprising, an isolated antibody, identifying susceptibility to a lipid disorder which comprises comparing the nucleotide sequence, where the difference between the suspected allele and the wild-type sequence identifies a sequence variation of FCHLI nucleotide sequence and a pharmaceutical composition. Also disclosed is a transgenic animal which carries an altered HYPLIP1 or FCHLI allele and a method for screening carries an altered HYPLIP1 or FCHLI allele and a method for screening carries an altered HYPLIP1 or FCHLI allele and a method for screening antibidies for inhibition or restoration of FCHLI, gene function as an antilipid disorder as a sequence with expression of FCHLI, for diagnosis or prognosis of predisposition to lipid disorder, and cancer and for treating a lipid disorder such as familial combined hyperlipidaemia, coronary artery disease, atherogenic lipoprotein phenotype, hyperapobetalipoproteinaemia, hypertriglyceridaemia, low density lipoprotein (LDL) subclass B, familial dyslipidemic hypertension, synthesis and cancer. The sequence presented is a PCR primer which was used to amplify part of the mouse HYPLIP1 locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rat; soluble adenylyl cyclase; sAC; glucose regulation; insulin release; pancreatic islet cell; cAMP production; physiological bicarbonate level; reverse transcriptase; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid molecule encoding soluble adenylyl cyclase polypeptide, useful for regulating insulin secretion from pancreatic islet cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
  acids of the encode polypeptides, where the 4 contiguous amino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 0.4%; Score 15.2; DB 1; Length 20; Local Similarity 85.0%; Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Soluble adenylyl cyclase (sAC) RT-PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTTGCATGCCTGCAGGTCGA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAY-1999;
26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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  amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1337
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8688888888888888888888888888888888
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The invention describes an isolated nucleic acid molecule (1), comprising a sequence encoding a polypeptide that comprises adenylyl cyclase catalytic domains C1 and C2 and rading at least 85% identity to a 50 kD-terminal domain comprising catalytic domains C1 and C2 of soluble

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adenylyl cyclase. The soluble adenylyl cyclase (sAC) comprises a fully defined sequence (S1) of 1608 or 1610 amino acids as given in the specification. (I) is useful for encoding soluble adenylyl cyclase. SAC is useful for glucose regulation of insulin release from pancreatic islet cells, and for regulating cAMP production. SAC is also useful as a sensor lin its capacity as an enzyme to determine physiological bicarbonate levels. This sequence represents a reverse transcriptase PCR primer used to analyse the expression of soluble adenylyl cyclase (SAC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to human orthologue of Aristaless homeobox gene, ARX. ARX gene is associated with infantile spasms (IS), non-specific X-linked mental retardation, X-linked myoclonic epilepsy and Partington syndrome. ARX gene is useful for diagnosing ARX-related disorders e.g., mild mental retardation, infantile spasms or dystonia. The present sequence is a PCR primer used for amplifying human ARX gene 3'
                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New ARX gene, useful for diagnosing ARX-related disorders e.g., mild mental retardation, infantile spasms or dystonia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homeobox gene; ARX; ARX-related disorder; X-linked myoclonic epileps
infantile spasm; mental retardation; Partington syndrome; diagnosis;
dystonia; human; untranslated region; UTR; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human ARX gene 3' untranslated region (UTR) amplifying primer, {
m F.}
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                                                                                                                                                                                                               0.4%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 1.5e+03;
rative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ative 0; Mismatches 3; Indels
                                                                                                                                                                             Sequence 20 BP; 4 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 25; 74pp; English.
                                                                                                                                                                                                                                                                                             1998 CAAGCAGCTGGTGGAGGACC 2017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1836 CTTCACGCTGGGGGCTCCC 1855
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                                                                                                                                                                                                                                                                                                                                                                                                                              AAL61164 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-SEP-2003 (first entry)
                                                                                                                                                                                                                   Query Match 0.4
Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           untranslated region (UTR)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-505184/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gecz J, Stromme P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 17; Conserv
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Matches
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Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2. useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
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Manoharan M;
                                                                                                                                                                                     antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss; 2'-methoxyethoxy-residue; phosphorothioate backbone.
                                                                                                                                                   Human mdm2 antisense oligonucleotide #145.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 147; 289pp; English.
                                     ADD21582 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                      02-DEC-2002; 2002WO-US038281.
                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2001; 2001US-00005344.
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                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 85.09
Watches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-577263/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mdm2 expression.
                                                                                                                                                                                                                                                                                                                             WO2003048315-A2.
                                                                                                                                                                                                                                                                                           sapiens.
                                                                                                               15-JAN-2004
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                                                                           ADD21582;
RESULT 1340
ADD21582/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1341
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes an isolated polypeptide (I) comprising a variant form of a mouse HYPLIPI polypeptide sequence (S1) or a human FCHLI.

Colypeptide sequence (S2), not given in the specification, where the variant form is associated with cancer, or an amino acid sequence having at least 65 % sequence identify to (S1) or (S2). A composition comprising or least 65 % sequence identify to (S1) or (S2). A composition comprising (I) is useful for treating or preventing cancer associated with expression of FCHLI. FCHLI gene or HYPLIPI gene and its product are useful for the study of metabolic pathway and cellular mechanism to identify other genes, receptors and relationships that contribute to inpid disorder and cancer. FCHLI gene or its fragments are useful in gene the treatment of lipid disorder or cancerous cells. The sequence to variation of FCHLI gene or HYPLIPI gene is also useful in the diagnosis and prognosis of predisposition to lipid disorder and cancer. Antisense polymucheotide sequences are useful in preventing or diminishing the expression of HYPLIPI or FCHLI locus. This sequence represents a primer
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                                                                                                                                                                                     cytostatic, antilipemic, gene therapy, peptide therapy, HYPLIPI, FCHLI, cancer, metabolic pathway, cellular mechanism, lipid disorder, familial combined hyperlipidaemia, mouse, PCR, primer, ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human FCHLI or mouse HYPLIPI polypeptide, useful for drug screening, peptide therapy of lipid disorder or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chatterjee A, Jong PD,
Wu C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used in the analysis of the mouse HYPLIP1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTTGCTTGCCTGCAGGCCCA 3670
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Ohmen J, Ross D, Tafuri S,
                                     ADB95852 standard; DNA; 20 BP
                                                                                                                                                   Mouse HYPLIP1 PCR primer #230.
                                                                                                                                                                                                                                                                                                                                                                                  07-SEP-2001; 2001US-00949427.
                                                                                                                                                                                                                                                                                                                                                                                                                          08-SEP-2000; 2000US-0231322P.
                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            BODNAR J S.
CASTELLANI L W.
CHATTERJEE A.
JONG P D.
LUSIS A J.
OHMEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-695901/66.
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TAFURI S.
                                                                                                                                                                                                                                                                                                           US2003054418-A1.
                                                                                                             04-DEC-2003
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Best Local Si
Matches 17;
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                                                                         ADB95852;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ROSS/)
(TAFU/)
(WUCC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BODN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (JONG/)
(LUSI/)
(OHME/)
                                                                                                                                                                                                                                                                       Mus sp.
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RESULT 1339
ADB95852
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The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a phosphorothioate backbone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide; human; mdm2; hyperproliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mdm2 antisense oligonucleotide #207.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1346 CTGAGATGGAGATGATGAAG 1365
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03-JUL-2003
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Haematopoietic cell; tyrosine kinase; hyperproliferative disorder; cancer; therapy; inflammation; diabetes; viral infection; inflammation; tumour; cytostatic; virucide; antisense therapy; antisense; human; phosphorothicate backbone; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human haematopoietic cell tyrosine kinase antisense oligo ISIS #150763.
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0
hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss; 2'-methoxyethoxy-residue; phosphorothioate backbone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
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                                                                                                                                                                                                                                                                                                                                              Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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Manoharan M;
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/*tag=
                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
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Best Local Similarity
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                                                                                                                                WO2003048315-A2
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modified_base
                                                                                          Homo sapiens
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Synthetic.
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The invention relates to a compound targetted to a nucleic acid molecule encoding haematopoietic cell protein tyrosine kinase. The compound inhibits the expression of haematopoietic cell protein tyrosine kinase and it specifically hybridises with the nucleic acid molecule encoding the tyrosine kinase. The site on the nucleic acid molecule encoding the tyrosine kinase. The antisense compounds are useful for modulating the expression of an active antisense compounds are useful for modulating the expression of haematopoietic cell protein tyrosine kinase and treating diseases or conditions associated with the expression of the tyrosine kinase, such as the reprofilerative disorders (e.g. cancer), inflammation, diabetes or a viral infection. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in pathway. The present functions of various members of a biological pathway. The present sequence is human haematopoietic cell tyrosine kinase antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense oligonucleotides targeted to nucleic acids encoding hematopoietic cell protein tyrosine kinase, useful for diagnosing treating cancer (e.g. leukemia), inflammation, diabetes or viral
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                                                                                                            /mod_base= OTHER
/note= "2'-0-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "2'-0-methoxyethyl (2'-MOE) nucleotides"
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85.0%; Pred. No. 1.5e+03;
iive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; Page 26; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1679 ACTICGGGCTGGCCCGGGAC 1698
methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 ACTITGCCTGGCCCGGGTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF88448 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2001; 2001US-00007010.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-DEC-2001; 2001US-00007010.
                                          1..5
/*tag= b
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                                                                                                                                                                                                                               /*tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 85.0
nes 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Borchers AH, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-811000/76.
                                                                                                                                                                                                                                                                                                                                                                              US2003125275-A1
                                      modified base
                                                                                                                                                                                         modified base
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The invention relates to antisense compounds, compositions and methods for modulating the expression of Sonic hedgehog (SHH). The composition comprises antisense compounds targetted to SHH. The antisense compound is used to inhibit the expression of SHH in cells or tissues and to treat an animal having a disease or condition associated with SHH, such as cancer, an autolimume disease or an inflammatory disorder. It is also useful in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The antisense compounds are useful in antisense gene therapy. The present sequence is an antisense oligonucleotide targetted to human SHH DNA. This sequence is used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a // Arag= Arage // Arag= a phosphorothioate // Anote= "This oligonucleotide has a phosphorothioate backbone and 2-"methyoxyethyl (2'-MOE) wings at the 5'
                                                                                                                                                                                                                                                                                                                                                                                         New compound for inhibiting Sonic hedgehog (SHH) expression in cells tissues and for treating an animal having a disease or condition associated with SHH, such as cancer, an autoimmune disease, or an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FGF receptor 2 antisense oligonucleotide, ISIS143404, SEQ ID 98.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; Vulnerary; Gene Therapy; Antisense; fibroblast growth factor receptor 2; FGF receptor 2; hyperproliferative disorder; cancer; developmental disorder; wound healing; ss; phosphorothioate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
/*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl reidues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; SEQ ID NO 33; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2004 GCTGGTGGAGGACCTGGACC 2023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 geregreaageacereagee 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                 16-NOV-2001; 2001US-00001844.
                                                                                                                                                                                                          16-NOV-2001; 2001US-00001844.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADH63144 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammatory disorder.
                                                                                                                                                                                                                                               (BENN/) BENNETT C F. (COWS/) COWSERT L M.
                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-897023/82.
                                                                                    US2003105041-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                                                                                                                                                                                                                                                                                                             Bennett CF,
                                                                                                                        05-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADH63144;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1345
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  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel polynucleotide isolated and purified from a human gene having any one of 935 fully defined sequences as given in specification, or a sequence having a base substitution. The invention further relates to: an oligonucleotide containing single nucleotide polymorphisms; a PCR primer set chosen from the combination of two DNA fragments from any one of 1202 fully defined sequences as given in specification, a labelling probe containing the SNP containing oligo; and a microarray equipped with the SNP containing oligo. The isolated human can further the snP containing oligo. The isolated human can of the invention is useful for detecting the single nucleotide polymorphisms in human gene. The isolated human gene is also useful for diagnosis of disease and determination of side effect to a medical agent. The isolated human gene is also effective in detecting single nucleotide polymorphisms one of the PCR primers used in the single nucleotide polymorphism one of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                        Novel polynucleotide useful for detecting single nucleotide polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sonic hedgehog; SHH; cancer; autoimmune disease; inflammatory disorder; antisense gene therapy; human; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    backbone where all cytidine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human SHH specific antisense oligonucleotide, ISIS 104352.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 9 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /\text{note} = "2' - \text{methoxyethyl reidues}" 16..20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "Phosphorothioate backbon
residues are 5-methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                      Claim 2; SEQ ID NO 2031; 704pp; Japanese
                                                                                                                                                                                                          (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         921 CTTCTTCCTGTTCATCCTGG 940
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/mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG93079 standard; DNA; 20 BP
                                                                                                                        12-FEB-2002; 2002JP-00034717
                                                                                                                                                                 12-FEB-2002; 2002JP-00034717
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*tag=
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                                        JP2003235571-A.
                                                                                                                                                                                                                                                                                                                in human gene.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified base
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  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-MAR-2004
                                                                                 26-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG93079;
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Gaps

vivlemore401-10.rng

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The present invention relates to antisense oligonucleotides (ADH63077-ADH63154) targeted to fibroblast growth factor (FGF) receptor 2 coding sequences (ADH63049 and ADH63056), which specifically hybridize with and inhibit FGF receptor 2 expression. The antisense oligonucleotides are useful for treating or preventing diseases or conditions associated with FGF receptor 2 in an animal, e.g. hyperproliferative disorders (particularly cancer of the colon, lung, breast or skin), or developmental disorders. The antisense compound may also be used in wound healing. The antisense compounds are useful for diagnostics, therapeutics, prophylaxis, or as research reagents or kits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchollation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
and 3' ends, which are 5 nucleotides in length. Also all
cytidine residues are 5-methylcytidine"
                                                                                                                                                                                                                                                                                               New antisense oligonucleotides for modulating expression of genes encoding fibroblast growth factor receptor 2, useful for treating hyperproliferative (e.g. cancer of the colon, lung, breast or skin) or developmental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match

0.4%; Score 15.2; DB 1; Length 20;
Best Local Similarity 85.0%; Pred. No. 1.5e+03;
Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 7 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; SEQ ID NO 98; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1945 TACATGATCATGCGGGAGTG 1964
                                                                                                                                                                                                                               Cooper SR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 racardarcardacederg 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ85121 standard; DNA; 20 BP
                                                                                                                                                           14-SEP-2001; 2001US-00954556.
                                                                                                                         12-SEP-2002; 2002WO-US029149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                               Freier SM,
                                                                                                                                                                                                                                                                WPI; 2003-354582/33.
                                                     WO2003024987-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200285308-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-OCT-2002.
                                                                                        27-MAR-2003
                                                                                                                                                                                                                             Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ85121;
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intiation codon, coding region, 5' or 3' end genomic flanking regions, 6' incitions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention of the invention and antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antifilammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed of profice and prophylacide of this patent is not represented in the printed of the file and prophylacide of the prophylacide of the printed of the form o
                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antisethmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel pharmaceutical composition, which has a
                                                                                                                                                                                                                 respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                    Katz E, Pabalan J, Aguilar D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 5 A; 9 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                     Claim 15; SEQ ID NO 363; 872pp; English.
                                                    Sandrasagra A, K
,, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2160 CCCGGCCCCACCCAGCAGTG 2179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ91733 standard; DNA; 20 BP
(EPIG-) EPIGENESIS PHARM INC.
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                                                  Li Y, San
Tang L,
                                                                                                                                   WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
es 17; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200285308-A2
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                                                                                                                                                                                                                                                                      ubiquinone
                                               Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ91733;
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23-APR-2002; 2002WO-US013135 24-APR-2001; 2001US-0286137P

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or The invention relates to a novel pharmaceutical composition, which has a at ftp.wipo.int/pub/published pct sequences Disclosure; SEQ ID NO 6975; 872pp; English. Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; (EPIG-) EPIGENESIS PHARM INC. WPI; 2003-229219/22 ubiquinone. Miller S, Nyce JW,

first active agent comprising an oligomeclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 6' or regions within 2.10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an enast airway dysfunction and a second active agent comprising an has antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, receptor, producing bronchodilation, increasing levels of demosine receptor, producing bronchodilation, increasing levels of the subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO

Sequence 20 BP; 11 A; 0 C; 1 G; 8 T; 0 U; 0 Other;

Gaps ö Match 0.4%; Score 15.2; DB 1; Length 20; Local Similarity 85.0%; Pred. No. 1.5e+03; les 17; Conservative 0; Mismatches 3; Indels Query Match Best Local S Matches

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RESULT 1348

ABZ85521 standard; DNA; 20 BP

ABZ85521;

Human oligonucleotide sequence. (first entry) 17-0CT-2003

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human; antisense; lung dysfunction; nasal airway dysfunction;

Homo sapiens,

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D;

Pabalan J, Aguilar D; Katz E, Sandrasagra A, Ka L, Shahabuddin S; Li Y, Sar Tang L, Nyce JW,] Miller S,

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Claim 15; SEQ ID NO 763; 872pp; English.

first active agent comprising an oligonuclectide antisense to the intration codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-axon junctions, or regions within 2'-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of or intinflammation, lung allergies, or a respiratory effect of an antiinflammation, lung allergies, or a respiratory disease or condition. Once the sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences The invention relates to a novel pharmaceutical composition, which has a

Sequence 20 BP; 9 A; 1 C; 3 G; 7 T; 0 U; 0 Other;

Gaps . 0 Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03; 0; Mismatches 3; Indels Query Match 0.4%; Best Local Similarity 85.0%; Matches 17; Conservative (

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ABZ85506/c ID ABZ85506 standard; DNA; 20 BP. RESULT 1349

ABZ85506;

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

antiniflammatory steroid; ubiquinone; antiniflammatory; antiallergic; antiantlammatory steroid; ubiquinone; antiantlammatory; antiallammatory; antiants gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human; antisense; lung dysfunction; nasal airway dysfunction;

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 5 and 3 intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antialergic, antiasthmatic, hypotensive, tumnosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also contininflammatory steroid in a subject, for reducing levels of an antiinflammatory steroid in a subject, for reducing levels of adenosine or lung surfactant in a subject, for reducing levels of denosine receptor, producing bronchodilation, increasing levels of adenosine or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the printed specification, but was obtained in electronic format directly from WIPO at the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                       Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                   respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 20 BP; 5 A; 0 C; 11 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                    Claim 15; SEQ ID NO 748; 872pp; English.
                                                    Sandrasagra A, K
L, Shahabuddin S;
(EPIG-) EPIGENESIS PHARM INC.
                                                    Li Y, Sar
Tang L,
                                                                                                                                  WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                       ubiquinone.
                                                    Nyce JW, ]
Miller S,
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Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels 317 ACCCCACTCCCTCCATCTCC 336

20 ACCCTACTCCCTCCAATTCC 1 g

ABZ87924 standard; DNA; 20 BP 17-OCT-2003 (first entry) ABZ87924; RESULT 1350 AB287924 110 AB28 AX AB28 XX AB28 XX BB Huma XX Huma XX Huma XX AB1 XM AB1 XM

Human oligonucleotide sequence.

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiansthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human; antisense; lung dysfunction; nasal airway dysfunction;

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Katz E, Pabalan J, Aguilar D;

Aguilar D; Katz E, Pabalan J, Sandrasagra A, K ,, Shahabuddin S; Li Y, San Tang L, Nyce JW, Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(8) antisense to specific gene(8) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or . biquinone

Disclosure; SEQ ID NO 3166; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entitiflammatory steroid in a subject, for reducing levels of denosine receptor, producing bronchodilation, increasing levels of ubiquinone or ung surfactant in a subject's tissue, or treating bronchocinstriction, lung surfactant in a subject's tissue, or treating bronchosisticition, lung antilammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed of processing in the printed of participal parts of this patent is not represented in the printed processing in the printed of parts in the printed parts at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 9 A; 2 C; 8 G; 1 T; 0 U; 0 Other;

Gaps ö Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels

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Gaps

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ADA26867 standard; DNA; 20 ADA26867/

RESULT 1351

(first entry) 20-NOV-2003 ADA26867;

Human FLJ20297 reverse PCR primer #151.

neoplastic growth marker, drug screening; cancer; tumour; gastrointestinal; prostate; breast; colorectal; diagnostic imaging; drug targeting; human; cytostatic; reverse transcription-PCR; RT-PCR; growth; detection; prediction; Metastasis; neoplastic primer; ss

Homo sapiens

WO2003031930-A2.

17-APR-2003.

02-OCT-2002; 2002WO-US031247.

09-OCT-2001; 2001US-0327332P.

(UYJO) UNIV JOHNS HOPKINS

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18-SEP-2003
                                                                                                                                                                                                                                                                                                                      Monia BP,
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                                                                                                                                                                                                                                                             (GAAR/)
(NERO/)
                                                                                                                                                                                                                                                                                                                                                                                                                           disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1353
ABD21736/c
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                                                                                                                                                                                The invention relaters to methods for internitying regions on incoppasition arker protein in a pumman patient, especially for detecting or predicting metastasis. The methods involve determining whether a neoplastic growth marker protein is overexpressed, either by the use of an antibody opecific for the protein, or by the use of PCR or hybridisation to detect nucleic acids encoding the marker proteins. A set of neoplastic growth markers are disclosed (SAGE (serial analysis of gene expression) tags for these are given in ADA26759-ADA26796), with protein tyrosine phosphatase type IVA member 3 (also known as PRL-3) being a preferred neoplastic growth marker. The neoplastic growth markers are specifically expressed at a higher level in metastatic cancers, compared with advanced and early stage cancers and normal cells frow which the cancer is derived.

Overexpression of the neoplastic growth markers is taken as an indication of that the tissue has a propensity to metastasiss. The invention also encompasses methods for treating a patient with an advanced or metastatic cancers. The methods of the invention are useful for identifying regions of neoplastic growth, for detecting or predicting metastatic cancers. The invention is paricularly applicable to incorpance or metastatic cancers. The invention is paricularly applicable.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                     human
                                                                  Identifying regions of neoplastic growth in a human body, useful for detecting or predicting metastasis, comprises administering to the humar body an antibody or peptide that specifically binds to a protein marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= b
/mod_base= Other
/note= "Phosphorothioate linkages. All cytosines are 5-
methyl cytosines"
1..5
                                                                                                                                                                        neoplastic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gastrointestinal, prostate, breast or colorectal cancers. Antibodies which bind to the neoplastic growth marker proteins are additionally useful for diagnostic imaging and for targeting cytotoxic or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transcription-PCR (RT-PCR) primer used to study the upregulation of neoplastic growth marker genes in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chemotherapeutic drugs. The present sequence represents a reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense; p38 mitogen activated protein kinase; p38 MAPK;
inflammatory disease; autoimmune disease; rheumatoid arthritis;
heart disease; ss; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                       The invention relates to methods for identifying regions of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
             Bardelli A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human p38 MAPK antisense oligonucleotide #20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
               Saha S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1385 ACATCATCAACCTGCTGGGC 1404
                                                                                                                                            Example 2; Page 22; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 ACTACATGAACCTGCTGGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM34234 standard; DNA; 20 BP
             Vogelstein B, Kinzler KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                  neoplastic growth.
                                           WPI; 2003-393457/37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADM34234;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1352
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The invention relates to an antisense oligonucleotide 8-30 nucleobases in length targeted to the 5'-untranslated region, translational start site, translational termination region or 3'-untranslated region of a nucleic acid molecule encoding a p38 mitogen activated protein kinase (MAPK). Where the antisense compound inhibits the expression of the p38 MAPK. The antisense oligonucleotide is useful for inhibiting the expression of p38 MAPK. The actisense oligonucleotide is useful for inhibiting the expression of p38 MAPK in cells or tissues. It is also useful for treating an animal having a disease or condition associated with p38 MAPK, e.g. an inflammatory or an autoimmune disease (e.g. rheumatoid arthritis) or a heart disease. In addition, the compound is used for disanostics, prophylaxis, or as research respents a p38 MAPK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity, lung, cancer; surfactant depletion, antiallergic, antiinflammatory; antiasthmatic, analgesic, hypotensive, immunosuppressive, cytostatic, cystic fibrosis, beta-adrenergic agonist; respiratory disease, pulmonary vasoconstriction, respiratory distress syndrome, allergic rhinitis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense oligonucleotides for modulating p38 mitogen activated protein kinase (MAPK) expression, useful for diagnosing, preventing or treating diseases associated with p38 MAPK, e.g. inflammation or heart
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35.0%; Pred. No. 1.5e+03;
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                                                                                                                                    /*tag= c
/mod base= Other
/note= "2'-methoxyethoxy nucleotides"
                             /mod_base= Other
/note= "2'-methoxyethoxy nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 9 A; 8 C; 1 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mckay R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 22; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2321 GIGIGIGIGIGIGIGI 2340
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-APR-1999; 99US-00286904.
15-AUG-2000; 2000US-00640101.
                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-2002; 2002US-00238442
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Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                   16. .20
/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-898587/82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GAARDE W A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MONI/) MONIA B P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MCKA/) MCKAY R.
                                                                                                       modified base
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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating controlled to respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The colligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression or cancer and can be anti-sense to the corresponding mann. CC The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention as antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mann or comprises of composition and/or surfactant hypoproduction are associated inflammation, allergies and/or bronchocomstriction and/or lung allergies, asthma, impeded respiratory disease, pulmonary compliances, pain, cystic fibrosis, allergic rhinitis, pulmonary crannerlant, melloctic behavior the production are associated transcribion, emphysema, chance in the degradation of the carget pulmonary comprises of pain, cystic fibrosis, allergic rhinitis, pulmonary therefore on a pulmonary disease, pulmonary diseases, pulmonary comprises.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
  disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ဌ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                               Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                  Katz E, Pabalan J,
emphysema; chronic obstructive pulmonary disease pulmonary transplantation rejection; ss; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; SEQ ID NO 748; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                             Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                                                            23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                            24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                                (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-093058/08.
                                                                                                                           WO200285309-A2
                                                                            Homo sapiens.
                                                                                                                                                                             31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                       Miller S,
                                                                                                                                                                                                                                                                                                                                                                                  Nyce JW,
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Sequence 20 BP; 5 A; 0 C; 11 G; 4 T; 0 U; 0 Other;

Gaps ; 0.4%; Score 15.2; DB 1; Length 20; 15.0%; Pred. No. 1.5e+03; 3; Indels 0; Mismatches 82.08; 17; Conservative Best Local Similarity Query Match Matches

317 ACCCCACTCCCTCCATCTCC 336

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20 ACCCTACTCCCTCCAATTCC 1 ABD21351 standard; DNA; 20 BP. RESULT 1354 ABD21351 ID ABD2 XX

ABD21351;

(first entry) 29-JUL-2004

Human transglutaminase-derived oligo SEQ ID 363.

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antialergic; antialinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; s; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, L Miller S,

WPI; 2003-093058/08.

ဌ Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 363; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating to comprising oligonucleotides, effective for alleviating to comprising oligonucleotides, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, contract depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regularing expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises. (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antianflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or traating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to inflammation, allergies and/or bronchoonstriction and/or lung inflammation, allergies and/or bronchoonstriction and/or bronchoonstriction and or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, the reduced adenosine content of the anti-sense oligos corresponding to the reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into preduce that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to it.

Sequence 20 BP; 5 A; 9 C; 4 G; 2 T; 0 U; 0 Other;

Query Match

DB 1; Length 20; 0.4%; Score 15.2;

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antisthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies, asthma, impeded respiration, respiratory
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                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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                       3; Indels
                                                                                                                                                                                                                                                                                                                                       Human calmodulin 2-derived oligonucleotide SEQ ID 3166.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Katz E, Pabalan J,
      Pred. No. 1.5e+03;
                       0; Mismatches
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                                                              2160 CCCGGCCCCACCCAGCAGTG 2179
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Tang L, Shahabuddin S;
                                                                                           CACGACTCCACCCAGCAGTG 20
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  82.0%;
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                                                                                                                                                                                                              ABD24154 standard; DNA; 20
                                                                                                                                                                                                                                                                                              (first entry)
                       17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-093058/08.
Best Local Similarity
Matches 17; Conserv
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Miller S,
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hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; 88; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation (a) or (a) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c)
distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary
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nucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                                                                                               Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                         Sequence 20 BP; 9 A; 2 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA497002-derived oligonucleotide SEQ ID 6975.
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                                                                                                                                                                                                                                                                                                                       2275 AGACTCAGTGCAGATGGAGA 2294
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABD27963 standard; DNA; 20
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Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABD27963;
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analgesic, hypotensive, immunosuppressive and cytostatic activity, is beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic thinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
      carrier and for use of the kit. The composition
instructions for adding a carrier and for use of the kit. The compo of the invention has antiallergic, antiinflammatory, antiasthmatic,
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Sequence 20 BP; 11 A; 0 C; 1 G; 8 T; 0 U; 0 Other;

Gaps ö th 0.4%; Score 15.2; DB 1; Length 20; Similarity 85.0%; Pred. No. 1.5e+03; 17; Conservative 0; Mismatches 3; Indels Local Similarity Query Match Matches

2824 ATATATATATATATAT 2843 1 ATATGAAATATATATAT 20 ਨੇ

ABD21751 standard; DNA; 20 BP RESULT 1357 ABD2175:

ABD21751;

(first entry) 29-JUL-2004

Human stanniocalcin-derived oligo SEQ ID 763.

respiratory tract inflammation, adenosine sensitivity, lung; cancer; surfactant depletion; antialinersametory; antiation analogesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens.

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, 1 Miller S,

Katz E, Pabalan J, Aguilar D;

WPI; 2003-093058/08.

ţ Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating conditions.

Composition also deenosine sensitivity, levels of adenosine (A) or (A) receptors, reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, configuration or cancer and carbon a sene encoding or regulating expression of a target polypeptide associated with lung airway or lung dystunction or cancer and can be anti-sense to the corresponding mRNA.

Composition of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung constructions for adding a carrier and for use of the kit. The composition of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antialflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostratic activity, is a composition composition composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition composition and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory chitammation, allergies, asthma, impeded respiration, respiratory bulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cystic fibrosis, allergies and cystic fibros lung, brain, heart, kidney, etc, tissue environment and thereby, to Claim 15; SEQ ID NO 763; 763pp; English. prevent any unwanted effects due to it

Sequence 20 BP; 9 A; 1 C; 3 G; 7 T; 0 U; 0 Other;

Gaps ; 0 0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels 17; Conservative Query Match Best Local Similarity Matches 17; Conserv

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ADF42870 standard; DNA; 20

ВР

ADF42870;

(first entry) 12-FEB-2004 Human RET proto-oncogene PCR primer SEQ ID NO 4.

human; ovarian cyst, RET; tumour; anticancer; taxol; cisplatin; adriamycin; epirubicin; cyclophosphamide; camptothecin; cytostatic; gene therapy; proto-oncogene; ss; primer; PCR.

WPI; 2004-038435/04

Novel ovarian cyst marker substance containing RET gene product, useful

05-APR-2002; 2002JP-00104215. 05-APR-2002; 2002JP-00104215. (SASA/) SASAKI H. JP2003289877-A. Homo sapiens. 14-OCT-2003.

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RESULT 1361
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                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to method of illumination and a biochip illumination apparatus that has a respective divergent diffuser proximate to each line of optical fibre faces for coupling and diffusing laser tight evently through the opposing sides of the glass substrate to illuminate the bioarray carried by the glass substrate. The illumination source includes a low power non-collimated laser emitting diode (LED). The divergent diffuser is made of silicon. The apparatus is useful for Dioarray illumination in a biochip reader. The invention provides a superior signal to noise ratio as compared with conventional illumination byochip manufacturing.
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                                                                                          This invention describes a novel human ovarian cyst marker containing the RET proto-oncogene. The RET gene can be used in manufacturing a pharmaceutical for treating an ovarian cyst or for treating a tumour which also comprises an anticancer compound chosen from a taxol, a cisplatin, an adriamycin, an epirubicin, a cyclophosphamide and a anticancer activity and can be used in gene therapy. ADF42867-ADF42884 represent PCR primers used to amplify the human RET proto-oncogene described in the disclosure of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biochip illumination apparatus includes respective divergent diffuser proximate to each line of optical fiber faces for coupling and diffusing laser light evenly through opposing sides of glass substrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide probe #52 used in biochip manufacturing
                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 10 A; 6 C; 4 G; 0 T; 0 U; 0 Other;
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                                               Claim 11; SEQ ID NO 4; 22pp; Japanese.
  as target in diagnosing ovarian cyst.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1810 TCCTTTGGGGTCCTGCTCTG 1829
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Best Local Similarity 85.C
Matches 17; Conservative
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ADE78926
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AC ADF78926
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DT 26-FEB-;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biochip illumination apparatus includes respective divergent diffuser proximate to each line of optical fiber faces for coupling and diffusing laser light evenly through opposing sides of glass substrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               111umination system; biochip illumination; divergent diffuser;
optical fibre; laser light; glass substrate; bioarray;
laser emitting diode; LED; biochip reader; biochip manufacturing; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to method of illumination and a biochip
                                                                                                                            Gaps
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                                                                 Length 20;
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                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide probe #63 used in biochip manufacturing.
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BP; 7 A; 0 C; 7 G; 6 T; 0 U; 0 Other;
                                                       Score 15.2; DB 1;
Pred. No. 1.5e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 74; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1350 GATGGAGATGATGAGATGA 1369
                                                                                                                                                                                          1350 GATGGAGATGATGAAGATGA 1369
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                                                                                                                                                                                                                                                 1 GATGATGATGATGATGA
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                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                              Query Match 0.4%;
Best Local Similarity 85.0%;
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      siochip manufacturing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYCH-) UNIV CHICAGO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-008450/01.
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      Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF78948;
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15-MAY-2003; 2003WO-US015493.
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13-NOV-2002; 2002US-0426324P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH51031 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 85.0 nes 17; Conservative
                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                                             Graham MJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               are 5-methylcytidines.
                                                                                                                                                                                                                           WPI; 2004-022840/02.
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WO2003097662-A1
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                                  27-NOV-2003
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                                                                                                                                                                                             Crooke RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH51031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a compound targeted to a nucleic acid molecule encoding Notchl and which inhibits the expression of Notchl. The compound, composition and methods are useful for treating a disease or condition associated with Notchl, such as an autoimmune disorder, aberrant apoptosis or a hyperproliferative disorder, e.g. cancer. They are also useful in research and diagnostics for modulating the expression of Notchl. The present sequence represents a human Notchl antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding Notch1, useful for treating an autoimmune disease, aberrant apoptosis or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           apolipoprotein B; ApoB; antiarteriosclerotic; cardiant; antidiabetic; anorectic; lipid; cholestrol metabolism; atherosclerosis; diabetes Type 2; obesity; hyperlipidaemia; cardiovascular; gene therapy; antisense; 2-0-methoxyethy! gapmer; phosphorothioate backbone; 2'-MOB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2'-MOE gapmer antisense oligo targeted to human ApoB DNA 3 - SEQ ID 385.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 0.4%; Score 15.2; DB 1; Length 20; Local Similarity 85.0%; Pred. No. 1.5e+03; es 17; Conservative 0; Mismatches 3; Indels .
                                                                                                                                                        ss; human; Notchl; autoimmune disorder; aberrant apoptosis; hyperproliferative disorder; cancer; antisense.
                                                                                                                      Human Notch1 antisense oligonucleotide ISIS 226826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; SEQ ID NO 22; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                410 GCCTGGTCATGGAAAGCGTG 429
                                                                                                                                                                                                                                                                                                                                                                                                                                         Koller E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 GGCTGGTCAGGGAAATCGTG 1
ADG88866/c
ID ADG88866 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADH18396 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                   30-MAY-2002; 2002US-00160497
                                                                                                                                                                                                                                                                                                                                                                      30-MAY-2002; 2002US-00160497
                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                         Freier SM, Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-022077/02.
                                                                                                                                                                                                                                                                US2003224513-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotide
                                                                                                                                                                                                                               Homo sapiens.
                                                                                     11-MAR-2004
                                                                                                                                                                                                                                                                                                04-DEC-2003
                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; ss.
                                                  ADG88866;
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Homo sapiens

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                                                                                                                                                                               acid molecule encoding human apolipoprotein B (ApoB) which specifically hybridises with and inhibits the expression of human apolipoprotein B. The compound of the invention demonstrates antiarteriosclerotic, cardiant, antidiabetic and anorectic activities and may be useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis, diabetes Type 2, obesity, hyperlipidaemia or cardiovascular disease. Furthermore, the compound has gene therapy applications. The current sequence is that of the 2'-O-methoxyebthyl (2'-MOE) gapmer antisense oligo of the invention which has 2'-MOE 'wings', a phosphorothicate backbone throughout and in which all cytidine residues
                                                                                                                                                       The invention relates to a novel antisense compound targeted to a nucleic
New antisense compound, useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis, diabetes Type 2, obesity, hyperlipidemia or cardiovascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Keratinocyte growth factor receptor; receptor; Thymic epithelial cell; MTS24; T lymphocyte; immunosuppressive; cytostatic; anti-HIV; virucide; gene therapy; vaccine; antiallergic; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Keratinocyte growth factor receptor PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                          Claim 1; SEQ ID NO 385; 405pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2567 ACCACGGGACATCACAGGGT 2586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 ACCACCTGACATCACAGGTT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boyd RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gill JW,
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The present sequence is that of a PCR primer for keratinocyte growth factor receptor (KGFR). RT-PCR was used to examine gene transcripts of importance to thymus organogenesis in marine thymic epithelial cells (TEC) expressing the cell surface glycoprotein MTS24. Transcripts for KGFR were found in both MTS24+ and MTS24- TECs. The present invention provides progenitor TECs which give rise to the complete thymus confirmment including both cortical and medullary epithelial lineages. These progenitor cells are identified by cell surface expression of MTS24. The cells can be used to prevent or treat diseases which can be alleviated by increasing the number of T cells and/or altering the T cell population of a subject. New thymic tissue derived from MTS24+ TECs is able to uptake appropriate T cell precursors such as the marrow cells from the ablood and convert them in the new thymic tissue to both new T cells and dendritic cells. MTS24+ TECs can be used
                                                New isolated MTS24 thymic epithelial progenitor cell, useful for treating cancer, allergy, autoimmune disease, produces a MTS24 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to induce tolerance in a subject to a graft from a mismatched donor, to prevent or treat an autoimmune disease, to enhance an immune response to an antigen (especially to treat HIV infection following HAART treatment), to treat an allergy, and to prevent or treat cancer (e.g. following chemotherapy, radiation therapy or bone marrow transplantation). The MTS24 TRCS may be genetically modified e.g. to express a transgene encoding a polypeptide, double-stranded RNA, catalytic nucleic acid or antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.2; DB 1; Length 20;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                   Example 5; SEQ ID NO 4; 70pp; English
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Best Local Similarity 85.05
WPI; 2004-082076/08
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Gaps ö

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Rat PIM1 antisense oligonucleotide K18. ADI26621 standard; DNA; 20 BP 15-APR-2004 (first entry) AD126621; RESULT 1364

phosphorothioate, analgesic; uropathic; antipruritic; cytostatic; antinflammatory; antiasthmatic; antisense; inhibition; gene therapy; tactile allodynia; urinary incontinence; neurogenic bladder; pruritus; tumour; asthma; primer; ss. modulator; PIM kinase family; locked nucleic acid; LNA;

Rattus sp.

WO2003106681-A2

24-DEC-2003

12-JUN-2003; 2003WO-EP006158

14-JUN-2002; 2002DE-01026702

(CHEF) GRUENENTHAL GMBH.

Erdmann V; Gruenweller A, Altan O, Kurreck J, WPI; 2004-142780/14

This invention describes novel oligonuclectides or polynuclectide constructs which are used in pharmaceutical or diagnostic compositions.

The oligonuclectides are used for identifying modulators of pain, based on the ability of labelled oligonuclectides or polynuclectide constructs to bind to an RNA. The oliginuclectides can also be used to diagnose disease associated with altered expression of genes of the PIM Kinase family by measuring binding. The oligonuclectides have at least one modified ribose, phosphodiester and/or base component. Particularly many, of the nuclectide is a phosphorothicate. The polynuclectide construct comprises a ribozyme, DNA enzyme, vector (particularly for expression) or peptide nucleic acid, most preferably a hammerhead ribozyme or Type 8-17 DNA crazyme. It may be attached to a carrier (preferably the proteins tet, transportin or ferritin) and/or encapsulated in a liposome. The products of the invention have analgesic, uropathic, antipruritic, cytostatic, antinfammarcory and antiasthmatic activity and can be used for antisense and catalytic inhibition of FIM kinases and for antisense gene therapy. The oligonuclectides are useful for treating (i) pain, especially chronic, hear-induced or inflammatory pain, or tactile allodynia and (ii) chronic, hear-induced or inflammatory pain, or tactile allodynia and contact and catalytic inhibition of plan kinases and contactile allodynia and contactile allodynia and contactile and contactile allodynia and contactile and contactil ö chronic, heat-induced or inflammatory pain, or tactile allodynia and (ii) urinary incontinence, neurogenic bladder symptoms, pruritus, tumours and inflammation, especially PIMI-kinase associated inflammation such as asthma, or generally any PIMI-related disease symptoms. They can also be used to screen for analgesic agents and for diagnosis of diseases associated with expression of PIM family genes. ADI26552-ADI26627 represent antisense oligonucleotides described in the disclosure of the against PIM1 kinase, useful for treating, e, tumors and inflammation, by gene Gaps ö Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03; 3; Indels Sequence 20 BP; 6 A; 6 C; 7 G; 1 T; 0 U; 0 Other; 0; Mismatches 1809 GICCTITGGGGICCTGCICT 1828 pain, urinary incontinence, 20 Greecresesarecreerer 1 Disclosure; Fig 3; 37pp; German New oligonucleotides directed 0.4%; Local Similarity 85.0 es 17; Conservative Query Match therapy Matches g ઠ

BP ADI32188 standard; DNA; 20 (first entry) 22-APR-2004 ADI32188; RESULT 1365 ADI32188

Murine H2K promoter PCR primer #2.

antiapoptotic; Bcl-2; Raf; transgenic; anticancer; Bcl-x; Bcl-w, Bfl-1; Brag-1; Mcl-1; Al; C-Raf; B-Raf; A-Raf; SP-C promoter; HSK promoter; cytostatic; cancer; PCR; primer; 88.

WO2004001367-A2

31-DEC-2003

24-JUN-2003; 2003WO-DE002156

24-JUN-2002; 2002DE-01028187.

MEDI-) MEDINNOVA GES MEDIZINISCHE INNOVATIONEN

Rapp UR,

Sedlacek H;

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel test system that contains agents
required for determination of at least one antiapoptotic member of the
Bell2 family and at least one member of the Raf family. The invention
involves a transgenic non-human mammal, especially a rodent, whose cells,
in a specific tissue, constitutively express Bell2 and Raf family
constitutively and a method of screening for potential anticancer agents.
The agents are detection reagents specific for mRNA or protein of the Bell
constitutively and a method of screening for potential anticancer agents.
The agents are detection reagents specific for mRNA or protein of the Bell
camplification. Preferred agents for detecting proteins are antibodies.
Specified antiapoptotic agents are Bell2, Bell-W, Bfl-1, Brag-1,
Mcl-1 or Al, and specified Raf members are C. B. or A.Raf, all
optionally mutated. The transgenic animals contain genes for Bell2 and
Raf under control of constitutive promoters, e.g. the SP-C promoter for
Raf under control of constitutive promoters, e.g. the SP-C promoter for
Raf under control of constitutive promoters, e.g. the SP-C promoter for
Raf under control of constitutive promoters, e.g. the SP-C promoter for
Raf under control of constitutive promoters, e.g. the SP-C promoter for
Raf under control of constitutive promoters, e.g. the SP-C promoter for
The products of the invention have cytostatic activity. The system is
used (i) to diagnose cancer, or the risk of developing it, in humans or
animals, or (ii) to screen for substances that, in cells that express
both Bell-2 members and Raf proteins, reduce expression or function of Bell
cancers. The method provides an early indication of the risk of
developing cancer. This sequence represents a PCR primer used to make the
cancers the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       methoxyethyl nucleotides. All cytidine methylcytidine."
                             Test system for detecting Bcl-2 and Raf family proteins, useful for diagnosis of cancer, or risk, and in screening for anticancer agents, also transgenic animals and cells for the test.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human, 98; antisense; PTP1B; protein phosphatase 1B; PTPN1; phosphorothioate backbone; hyperproliferative condition; cancer; cytostatic; antidiabetic; anorectic; type 2 diabetes; obesity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "OTHER= phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense DNA oligo to target human PTP1B DNA SegID 305.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 11 C; 1 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1013 AGATCTCCCGCTTCCCGCTC 1032
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nucleobases are 5'
                                                                                        Example 1; Page 13; 22pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 ACATCTCCCGCATCCCACTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
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Best Local Similarity 85.0
Matches 17; Conservative
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WPI; 2004-071776/07
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modified_base
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This invention relates to novel compositions and methods for modulating the expression of PTP1B (also known as protein phosphatase 1B and PTPN1). Specifically, it refers to antisense compounds that can target and hybridise with a nucleic acid molecule encoding PTP1B, as well as splice variants thereof and inhibit expression accordingly. PTP1B is a tyrosine phosphatase that plays an essential regulatory role in signalling mediated by the insulin receptor and as such is useful for treating diseases such as type 2 diabetes and obesity. Furthermore, PTP1B can suppress transformation of oncogenic genes, such that compositions of this invention can also be used to treat hyperproliferative conditions including cancer. Accordingly, these compounds can be described as having cytostatic, antidiabetic and anorectic activities. This oligonucleotide sequence is an antisense DNA oligo that targets human PTP1B DNA, and which has a phosphorothicate backbone and 2'-O-methoxyethyl wings, used in an exemplification of the invention.
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/note= "OTHER= 2' methoxyethyl nucleotides. All cytidine nucleobases are 5' methylcytidine."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding PTP1B, useful for treating a disease/condition associated with PTP1B, such as cancer, diabetes or obesity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Butler MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Notch1 antisense oligonucleotide seq id 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; SEQ ID NO 305; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cowsert LM, Wyatt JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        416 TCATGGAAAGCGTGGTCCC 435
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                                                                                                                                                                                                                                                                                                                                            31-JUL-2000; 2000US-00629644.
14-MAY-2001; 2001US-00854883.
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nes 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
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/*tag=
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                                                                                                    US2003220282-A1
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bhanot S,
Freier SM;
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The present invention relates to a novel flower-bud formation suppressor gene. The invention is useful for regulating the time to flowering of a plant, conferring an early flowering property on a plant, or producing an early flowering plant, e.g. a perennial fruit tree. The present sequence is MÖTFL (Malus x domestica TERNINAL FLOWER) gene amplifying RACE (rapid amplification of cDNA ends) PCR primer.
                                                                                                                                                                                                                                                                 New flower-bud formation suppressor nucleic acid, useful for regulating the time to flowering of a plant, conferring an early flowering property on a plant, or producing an early flowering plant, e.g. a perennial fruit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New compound, particularly an antisense oligonucleotide targeted to a nucleic acid encoding KIAA1531, useful for treating cancer, chronic inflammation or conditions involving hyperactivation of angiogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; KIAA1531; hyperproliferative disorder; cancer; angiogenesis hyperactivation; chronic inflammation; ss; antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human KIAA1531 antisense oligonucleotide ISIS #208221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 4 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 15; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3126 AACTTATTGACAACCGAGAA 3145
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                                                                                                               20-JUN-2002; 2002JP-00180289.
                                                                           26-FEB-2003; 2003US-00372909
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                                                                                                                                                    (NAAG-) NAT AGRIC RES
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                                                                                                                                                                                          Kotoda N, Wada M,
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US2003237109-A1
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                                    25-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADJ46797;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes an antisense compound, having a sequence comprising 8-50 bp targeted to a nucleic acid encoding Notchl. The compound specifically hybridises with the nucleic acid encoding Notchl and inhibits its expression. The compound has cytostatic and immunosuppressive properties and is suitable for used in gene therapy. The compound is useful for preparing a composition for treating a disease or condition associated with Notchl e.g., developmental, autoimmune or hyperproliferative disorders, such as cancer. This sequence represents an antisense oligonucleotide specific to human Notchl.
                                                                                                                                                                        /mod_base= OTHER
/note= "OTHER= 2'-O-methyoayethyl substituted nucleoside
amidites"
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                                                                                                               /note= "OTHER= 2'-O-methyoayethyl substituted nucleoside
                  /note= "OTHER= Phosphorothioate backbone, all cytidine
residues are methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding Norchl, useful for preparing a composition for treating a disease associated with Notchl e.g. autoimmune or hyperproliferative disorders, such as cancer.
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85.0%; Pred. No. 1.5e+03;
cive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 22; 91pp; English.
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                                                                        /*tag= a
/mod_base= OTHER
/mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                       30-MAY-2002; 2002US-00160497
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                                                                                                                                                                                                                                                                                                                                                                                                                           (FREI/) FREIER S M.
(DOBI/) DOBIE K W.
(KOLL/) KOLLER E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-042171/04
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                                                                                                                                                                                                                                                                       US2003225019-A1
                                                        modified base
                                                                                                                                                      modified base
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                      The invention relates to a compound targeted to and which specifically hybridises with a nucleic acid molecule encoding KIAA1531 and inhibits the expression of KIAA1531. The compound, composition and methods are useful for treating a disease or condition associated with KIAA1531, such as a hyperproliferative disorder, e.g. cancer, a disease or condition involving hyperactivation of angiogenesis, or chronic inflammation. They are also useful in research and diagnostics for modulating the expression of KIAA1531. The present sequence represents a human KIAA1531 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for treating a disease or condition associated with KIAA1531, such as a hyperproliferative disorder, e.g. cancer, a disease or condition involving hyperactivation of angiogenesis, or chronic inflammation. They are also useful in research and diagnostics for modulating the expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a compound targeted to and which specifically hybridises with a nucleic acid molecule encoding KIAA1531 and inhibits the expression of KIAA1531. The compound, composition and methods are
                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compound, particularly an antisense oligonuclectide targeted to a nucleic acid encoding KIRA1531, useful for treating cancer, chronic inflammation or conditions involving hyperactivation of angiogenesis.
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                                                                                                                                                                                    0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; KIAA1531; hyperproliferative disorder,; cancer; anglogenesis hyperactivation; chronic inflammation; ss.
                                                                                                                                                              Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                          Human KIAA1531 target sequence ISIS #125855.
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Example 15; SEQ ID NO 74; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dobie KW;
                                                                                                                                                                                                                                              1610 AGTGCATCCACAGGGACCTG 1629
                                                                                                                                                                                                                                                                      AGTTCACCCACAGGCACCTG 20
                                                                                                                                                                                                                                                                                                                                         ADJ46851 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                    17;
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                                                                                                                                                                                                                                                                                                                                                                    ADJ46851;
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Matches
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The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                             human; single nucleotide polymorphism; SNP; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 0 A; 11 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 6044; 2627pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleotide polymorphism in human gene.
1610 AGTGCATCCACAGGGACCTG 1629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     175 GACGAAGACGGGGAGGACGA 194
                                    Н
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                                      20 AGTICACCCACAGGCACCIG
                                                                                                                                                                                                                                                                    Primer of the invention #2735.
                                                                                                                                              ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-MAR-2002; 2002JP-00064373.
                                                                                                                                              ADK97015 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK97839 standard; DNA; 20
                                                                                                                                                                                                                           (first entry)
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Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-093977/10.
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                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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                                                                                                                                                                                    ADK97015;
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                                                                                                     RESULT 1371
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                                                                                                                        ADK97015,
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Gaps

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Gaps ö

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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LRH1 (also known as NR5A2). LRH1 is a monomeric orphan nuclear receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside linkage, a phosphorothioate linkage; at least one modified sugar moiety,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                             phosphorothicate; 2' MOE; breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis; hepatocellular carcinoma; aromatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antiinflammatory; virucidal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                methoxyethyl (2' MOE) nucleotides. All ses are 5-methylcytidine."
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                                                                                                                                                                                                                                                                                                                    Antisense DNA oligo used to modulate human LRH1 expression SeqID 1221.
                                                                                                                                                                                                                                                                                                                                                         human; 88; liver related homologue-1; LRH1; NR5A2; antisense;
    Indels
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/label= OTHER= phosphorothioate backbone
    3;
    0; Mismatches
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/note= "OTHER= 2' met
cytidine nucleobases
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                                           2314 GGTCTGTGTGTGTGTGTG 2333
                                                                                   GGGAGGTGTGTGTGTGTG 1
                                                                                                                                                                                       ADJ16671 standard; DNA; 20 BP
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  17;
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                                                                                                                                                                                                                                 ADJ16671;
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                                                                                                                                                RESULT 1374
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    Matches
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                                                                                                                                                                                                                                                                                          The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single nucleotide polymorphism in human gene.
                                                                                                                                                                 polynucleotide useful for PCR amplification along with two DNA ent from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 20 BP; 2 A; 2 C; 8 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                  Claim 2; SEQ ID NO 6868; 2627pp; Japanese
                                                                                (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
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                                                                                                                                                                                                             nucleotide polymorphism in human gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2320 TGTGTGTGTGTGCGTGTG 2339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%;
08-MAR-2002; 2002JP-00064373.
                                       08-MAR-2002; 2002JP-00064373.
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ADK97714 standard; DNA; 20
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Best Local Similarity
                                                                                                                          WPI; 2004-093977/10.
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Best Local Similarity
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06-MAY-2004

ADK97714;

16-SEP-2003

Synthetic

17;

Matches

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RESULT 1373

ADK977

fragment

Novel

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a 2'-O-methoxyethyl (2' MOE) and at least one modified mucleobase, a 5-methylcytidine. These antisense compounds are useful for treating or diagnoshing a disease associated with LRHI, such as breast cancer, dyslipidaemia, atherosclerosis, low HDE (high density lipoprotein), high LDE (low density lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit cytostatic, antilipaemic, antiarteriosclerotic, anorectic, hepatotropic, litholytic, antilnflammatory and virucidal activities. This oligouscleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRHI protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human; ss; liver related homologue-1; LRH1; NR5A2; antisense; bobsphorothicate; 2' MDS; breast cancer; dyelipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hippercholesterolaemia; gall scone; triglyceridaemia; obesity; hepatitis; hepatcellular carcinoma; aromatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatcorropic; litholytic; antiliflammatory; virucidal.
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cytidine nucleobases are 5-methylcytidine."
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                                                                                                                                                                                                                                                   0.4%; Score 15.2; DB 1; Length 20; 35.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                    3; Indels
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                                                                                                                                                                                                                    Sequence 20 BP; 7 A; 7 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                    0; Mismatches
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                                                                                                                                                                                                                                                                                                                 2306 AGAGCTTTGGTCTGTGTG 2325
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/note= "OTHER= 2'
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/note= "OTHER= 2
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nes 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1375
                                                                                                                                                                                                                                                                                    Matches
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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRHI) and splice variants thereof. Specifically, it refers to compositions 8-10 nucleobases in clength that target a portion of an active site on the nucleic acid molecule encoding LRHI (also known as NR5A2). LRHI is a monomeric orphan cuclear receptor protein that functions as a tissue specific or transcription factor. The present invention describes antisense or inkage, a phosphorothoate linkage, at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MOE) and at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MOE) and at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MOE) and at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MOE) and at least one modified sugar moiety, a disanse associated with LRHI, such as beast cancer, dyalipidaemia, atherosclerosis, low HDL (high density lipoprotein), high complement, observed acute or chronic chaptitis, as well as hepaticellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit containfilammatory and virucidal activities. This oligonucleotide sequence is an antisense DNA oligo used to modulate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
                                        New antisense oligonucleotides targeted to a nucleic acid encoding liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; ss; liver related homologue-1; LRH1; NR5A2; antisense; hospbrorchhiotete, 2' MOSE; breast cancer; dyalipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis; antilipaemic; arctinoma; aromatese; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antiinflammatory; virucidal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                            related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; Live 0; Mismatches 3; Indels
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/label= OTHER= phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression of the human LRH1 protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 12 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                    Example 15; SEQ ID NO 1999; 909pp; English.
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/*tag= a
/mod_base=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17; Conservative
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Best Local Similarity
WPI; 2004-083054/08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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ADL00873 standard; DNA; 20
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nes 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                            WO2004009541-A2
                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-MAY-2004
                                                                                                                                                                                                                                                                                                                              29-JAN-2004
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bhat BG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1378
ADL00873/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the expression of liver related homologue-1 (LRH1) and splice variants the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in clearly that target a portion of an active site on the nucleac acid molecule encoding LRH1 (also known as NESA2). LRH1 is a monomeric orphan conclear receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified internucleoside oligonucleotides that comprise at least one modified internucleoside at linkage, a phosphorothicate linkage; at least one modified sugar moiety, and agnosing a disease associated with LRH1, such as breast cancer, disgnosing a disease associated with LRH1, such as breast cancer, disgnosing a disease associated with LRH1, such as breast cancer, disgnosing a disease associated with LRH1, such as breast cancer, thigh density lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B vius-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit cyclostatic, antilibermic, antil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to novel antisense compounds useful for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antilipaemic, Cardiovascular; Analgesic; Antianginal; Antisense therapy; Human; Endothelial Lipase; dyslipidaemia; high density lipoprotein; HDL; cardiovascular disorder; metabolic syndrome X; ss.
                                               /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligo used to modulate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression of the human LRH1 protein of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide sequence is an antisense DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; SEQ ID NO 1052; 909pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGCACAGCTACCTTTAAA 20
                          OTHER
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                                                                                                                                                                                                                          01-JUL-2003; 2003WO-US020865
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                            /mod_base=
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Les 17; Conservative
                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
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                                                                                                                                                                           08-JAN-2004
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Homo sapiens

RESULT 1377

ADJ21727

Matches

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                                                                                 /*tag= a //*tag= //*tag= a //*tag= //*tag= //*tag= //*nd base= OTHER //*note= This oligonucleotide has a phosphorothioate backben and 2-'methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 4 nucleotides in length. Also all and 3' ends, which are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; VEGF co-regulated chemokine-1; VCC-1; vascular endothelial growth factor; ss; antisense compound; vascular endothelial growth factor; ss; antisense compound; phosphorothicate linkage; 2'-0-methoxyethy! sugar moiety; beathylcytosine; antisense oligonucleotide; diabetes; immunological disorder; cardiovascular disorder; neurological disorder; ischaemia; reperfusion injury; cancer; angiogenic disorder; haemangioma; tumour angiogenesis; rheumatoid arthritis; atherosclerosis; psoriasis; fibrosis; myocardial infarction; wound healing; bone fracture; cartilage damage; tissue regeneration; organ regeneration; periodontal disease; gut regeneration; atrial fibrillation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to antisense oligonucleotides (ADJ21603-ADJ25510) targeted to human Endothelial Lipase (EL) coding sequence (ADJ25517), where the antisense oligonucleotide specifically hybridises with and inhibits the expression of EL. The antisense oligonucleotides are useful for modulating the expression of endothelial lipase in cells or tissues to treat diseases associated with EL expression, such as dyslipidaemia, low high density lipoprotein (HDL), cardiovascular disorder or metabolic syndrome X. In addition, the oligonucleotides are used for diagnostics, prophylaxis, or as research reagents or kits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human VEGF co-regulated chemokine-1' DNA antisense oligonucleotide #406.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; SEQ ID NO 125; 1007pp; English
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1842 GCTGGGGGCTCCCCGTACC 1861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 GCTGCGGGGAGCCCCGTACC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-JUL-2003; 2003WO-US022410.
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/mod_base= OTHER /note= "OTHER phosphorothicate backbone. All cytidine residues are 5-methylcytidines"

Location/Qualifiers

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1. .20 /*tag=

/mod_base= OTHER /note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"

/mod_base= OTHER /note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"

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/*tag=

16. .20

l. .5 /*tag=

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gene therapy; endothelial specific molecule-1; ESM-1; ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-AUG-2003; 2003WO-US025833.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-AUG-2002; 2002US-0404495P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Griggs DW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Weinstein EJ,
                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                         modified base
                                                                                                                                                                                                                                                            modified base
                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
        The invention relates to an antisense compound targeted to a nucleic acid molecule encoding human vascular endothelial growth factor (VEGF) co-
cegulated chemokine-1 (VCC-1), and which specifically hybridises with and inhibits the expression of VCC-1. The invention also relates to a composition comprising the antisense compound, a method of inhibiting the expression of VCC-1. The invention also contacting the calls or tissues compound and a method of treating a human cartisgues with the antisense compound and a method of treating a human cartisgues with the antisense compound to an animal to inhibit expression of VCC-1. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothicate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one modified sugar moiety, preferably a 2'-0-comprises at least one condition. The antisense oligonucleotide preferably is a chimeric oligonucleotide. The antisense compound is useful for treating a disease or condition associated with VCC-1, such as diabetes, an immunological disorder, a cardiovascular disorder, a neurological disorder, a cardiovascular disorder, a neurological disorder, a perfusion injury, cancer or an angiogenic attended and sorder, e-g. haemangioma, tumour angiogenesis, rheumangiod arthritis, attended of bone fractures and cartilage damage, for regeneration of tissues or organs, for treating periodonal diseases, for gut protection or regeneration, for treatment of lung or liver fibrosis an antisense or organs, for treatment of lung or liver fibrosis an antisense management of artial fibrillation. This sequence regeneration for the antise of the condition of the conditio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide targeted to DNA encoding the human VCC-1 polypeptide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                  New antisense compounds targeted to a nucleic acid molecule encoding vascular endothelial growth factor co-regulated chemokine-1 (VCC-1), useful for treating VCC-1-associated disorders, e.g. diabetes or a neurologic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 406; 336pp; English.
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                                                                                                                       19-AUG-2003; 2003WO-US025891
                                                                                                                                                            19-AUG-2002; 2002US-040484P
                                                                                                                                                                                                 (PHAA ) PHARMACIA CORP
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                                          WO2004016224-A2.
                                                                                                                                                                                                                                         Weinstein EJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the invention
        Homo sapiens.
                                                                                   26-FEB-2004
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                                                                         comprising 8-30 bp targeted to a nucleic acid encoding endochelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                       The invention describes a new antisense compound, having a sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 15.2; DB 1; Length 20; 35.0%; Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 10 A; 1 C; 0 G; 9 T; 0 U; 0 Other;
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Claim 3; SEQ ID NO 1749; 555pp; English
                                                                                                                                                                                                                                                                                                                                       endothelial specific molecule-1 (ESM-1)
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cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective;

Human ESM-1 antisense oligonucleotide segid 1749.

(first entry)

03-JUN-2004

ADL59500;

ADL59500 standard; DNA; 20 BP.

vivlemore401-10.rng

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mFGES-1; mFGES-1; inhibitor; microsomal prostaglandin E2 synthase; mFGES-1; mFGES-1; inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:174.

01-JUL-2004 (first entry)

ADM13987;

/mod_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

Location/Qualifiers

Key modified_base

Homo sapiens

Synthetic

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*tag=

/*tag= c /mod_base= OTHER /note= "2'-O-methoxyethyls"

25-SEP-2003; 2003WO-US030374. 25-SEP-2002; 2002US-0413549P

WO2004028458-A2

(PHAA) PHARMACIA CORP.

WPI; 2004-305094/28.

Gierse JK;

/mod_base= OTHER /note= "2'-0-methocyethyls"

16. .20

modified base

1..5 /*tag= a

modified base

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The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
                                           cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1; Banofer; diabetes; cancer; ischaemia; reperfusion injury; anglogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing a composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                                                                                                                                         /note= "OTHER= phosphorothioate backbone. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                    /*tag= c
/mod_base= OTHER
/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 12 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                Human ESM-1 antisense oligonucleotide seqid 1562.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            endothelial specific molecule-1 (ESM-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; SEQ ID NO 1562; 555pp; English.
                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                             /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-AUG-2003; 2003WO-US025833.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-AUG-2002; 2002US-0404495P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Weinstein EJ, Griggs DW;
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modified_base
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                                                                                                                                                         Homo sapiens
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid mpGES-1 in chibits its expression; (2) a method of inhibiting the expression of mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and mpGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiavascular activities, and can ophthalmological, immunomodulatory and cardiavascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Aliahime's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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Gaps

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/ Match 0.4%; Score 15.2; DB 1; Length 20; Local Similarity 85.0%; Pred. No. 1.5e+03; No 17; Conservative 0; Mismatches 3; Indels

Query Match Best Local Si Matches 17,

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ADM13987 standard; DNA; 20

RESULT 1381 ADM13987

SXS

RESULT 1382

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human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8.30 by targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal mpGES-1 in cells or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antistlammatory, neuroprotective, antiatrhitic, vasotropic, antidiabetic, immunomodulator, cardiant, neuroprotective, sactionities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 8 A; 8 C; 3 G; 1 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "2'-0-methocyethyls"
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/note= "2'-O-methoxyethyls"
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modified_base
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                                                                                                                                          Gaps
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ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                            Length 20;
                                                                                                                                       3; Indels
                                           Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                         Score 15.2; DB 1;
Pred. No. 1.5e+03;
                                                                                         Query Match
0.4%; Score 15.2; D
Best Local Similarity 85.0%; Pred. No. 1.5e
Matches 17; Conservative 0; Mismatches
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/note= "2'-O-methocyethyls"
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                     1612 TGCATCCACAGGGACCTGGC 1631
                                                                                                                                                                                                                ||| |||||||| || ||||||| TGCTTCCACAGAGAACTGGC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                   ADM15219 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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ADMISSI 11D ADMISSI ADMISSI

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Gaps

residues are 5-methylcytidines"

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38-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate, human; microsomal prostaglandin E2 synthase; mPGBS-1; mPGBS-1; inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiavascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; acadiovascular disorder; neurological disorder; se.
                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 8 A; 8 C; 3 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                       Claim 4; SEQ ID NO 1350; 132pp; English.
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25-SEP-2002; 2002US-0413549P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17; Conservative
                          (PHAA ) PHARMACIA CORP
                                                                              WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                    Gierse JK;
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                                                                                                                                               ischemia.
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Matches
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromsome 9, more specifically to pay14.3. The present invention also describes! (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding (mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues, and dishabetic acid encoding having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiant, neuroprotective, copthalmological, immunomodulatory and cardiant activities, and can can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation Alzheimer's
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                                                                                                                                                                                                                                /mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                                     /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. 20
/*tag= c
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Best Local Similarity
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modified_base
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cytostatic; antidiabetic;
        immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                          /note= "phosphorothioate linkages and all cytidine
 microsomal prostaglandin E2 synthase inhibitor;
                                                                                                                                                                     residues are 5-methylcytidines
                                                                                                                                                                                             /mod_base= OTHER
/note= "2'-0-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to old an integrated to a nucleic acid encoding maying a sequence comprising 8-30 bp targeted to a nucleic acid mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (3) a method of inhibiting the expression of mpGES-1 in cells or tissues; and anisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiavascular activities, and can confit in a mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
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Claim 4; SEQ ID NO 1641; 132pp; English.
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Claim 4; SEQ ID NO 1312; 132pp; English.
WPI; 2004-305094/28.
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic, antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antidiammatory; neuroprotective; onotropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for treating e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
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/mod_base= OTHER
/note= "2'-0-methocyethyls"
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                                                                                                      ADM15125 standard; DNA; 20
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Query Match

Best Local Similarity 85.0
Matches 17; Conservative

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RESULT 1388
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           antinfilammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer: disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; acrdiovascular disorder; neurological disorder; se.
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                                                                                                                                                                                                                                            0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
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antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                               Sequence 20 BP; 2 A; 12 C; 3 G; 3 T; 0 U; 0 Other;
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/note= "2'-0-methocyethyls"
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/mod_base= OTHER
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Best Local Similarity 85.0°
Matches 17; Conservative
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mFGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to gene is located on chromosome 9, more specifically to organize mpGES-1 and antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibite its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antianflammatory, neuroprotective, nootropic, antianflammatory, neuroprotective, nootropic, antianflammatory, neuroprotective, and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or distinct associated with mPGES-1 e-g., inflammation, all and an antisense compound and antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e-g., inflammation, Alzheimer's
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e.g.,
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/note = "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
encoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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/mod_base= OTHER
/note= "2'-0-methoxyethyls"
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16. .20
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Matches 17; Conservative
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, ophthalmological, immunomodulatory and cardiavascular activities, and can confitted as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or discontine associated with mpGES-1 e.g., inflammation, Alzheimer's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 1399; 132pp; English.
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                                                                                                                           25-SEP-2003; 2003WO-US030374
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les 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mEGES-1). The human mEGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mEGES-1 and inhibite its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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/note= "phosphorothioate linkages and all cytidine
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85.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 3; Indels
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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Location/Qualifiers
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Best Local Similarity
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ADM15498

Homo sapiens

Gaps

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Indels

0; Mismatches

2326 TGTGTGTGTGTGTGTGTG 2345

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Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03;

0.4%;

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mmcGES-1 gene is located on chromosome 9, more specifically to office on the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding medGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal acidsase or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamaticity, neuroprotective, noctropic, antidiamaticy, namunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound to pe used as mPGES-1 inhibitors and in gene therapy. The antisense compound condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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/mod_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

Location/Qualifiers

sapiens

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/mod_base= OTHER /note= "2'-0-methocyethyls" 16. .20

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/note= "2'-0-methoxyethyls"

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; imPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antitinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1138
Sequence 20 BP; 10 A; 8 C; 0 G; 2 T; 0 U; 0 Other;
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                           Query Match
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Matches 17; Conservative
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                                                                               chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; nardiavosacular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
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                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1685.
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/note= "2'-0-methocyethyls"
16. .20
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New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                            25-SEP-2003; 2003WO-US030374.
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WO2004028458-A2
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                having a sequence comprising 8-30 bp targeted to a nucleic acid encoding medESS-1, which specifically hybridise with the nucleic acid meDESS-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, archimit, neuroprotective, antinfilammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological; immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammatory and activities and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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  9q34.3. The present invention also describes: (1) antisense compounds,
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Matches 17; Conservative
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to gene is located on chromosome 9, more specifically to gray a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal cantisense oligonucleotides and antisense one compounds have cytostatic, antidabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound to be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, a disease or condition associated with mPGES-1 e.g., inflammation in injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 1640; 132pp; English.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 3034.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, in cells or tissues; and (3) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, ophthalmological, immunomodulatory and cardiavascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation Alzheimer's
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neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; alzheimer. a diesase; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
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85.0%; Pred. No. 1.5e+03;
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/note= "2'-O-methocyethyls"
16. .20
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         '*tag= a
mod_base= OTHER
'note= "2'-0-methocyethyls"
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic,
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/mod_base= OTHER
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                                                                ADM14988 standard; DNA; 20 BP
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ADM14988/c
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ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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/note= "phosphorothioate linkages and all cytidine
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Best Local Similarity 85.0%;
Matches 17; Conservative
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mFGES-1 gene is located on chromosome 9, more specifically to oligan mPGES-1 gene is located on chromosome 9, more specifically to man mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and not mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antinflammatory, neuroprotective, notropic, antisathritic, vasotropic, ophthalmological, immunomodulatory and cardiant activitities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or confident activities and in gene therapy. Alzheimer's confident associated with mPGES-1 e-g., inflammation, Alzheimer's confident activities and action associated with mPGES-1 e-g., inflammation, Alzheimer's confident activities and can be used do the propertice of 
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                                                     Claim 4; SEQ ID NO 1665; 132pp; English.
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Matches 17; Conservative
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to human mPGES-1 gene is located on chromosome 9, more specifically to human mPGES-1 and sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cyrostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, contininflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Altheimer's condition associated with mPGES-1 e.g., inflammation, Altheimer's condition associated with mPGES-1 e.g., inflammation, Altheimer's condition associated with mPGES-1 e.g., inflammation, and antisense compound can appear to the condition associated with mPGES-1 e.g., inflammation, and antisense compounds the condition associated with mPGES-1 e.g., inflammation, and antisense compounds the condition associated with mPGES-1 e.g., inflammation, and antisense composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, and antisense composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, and antisense composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, and antisense composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, and antisense composition for treating and antisense comp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1431.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 10 A; 8 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 1505; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2326 TGTGTGTGTGTGTGTG 2345
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                                                                                   25-SEP-2003; 2003WO-US030374
                                                                                                                                              25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                      (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                        WPI; 2004-305094/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                    Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                schemia.
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ADM15244/c
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; imPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; sont inflammation; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1566.

/mod_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

Location/Qualifiers

Ω

1. .20 /*tag=

modified_base

Homo sapiens.

Synthetic.

/mod_base= OTHER /note= "2'-0-methocyethyls" 16. 20 /*tag= c

modified base

′*tag= a

modified base

/mod_base= OTHER /note= "2'-O-methoxyethyls"

25-SEP-2003; 2003WO-US030374. 25-SEP-2002; 2002US-0413549P.

WO2004028458-A2

08-APR-2004,

(PHAA) PHARMACIA CORP

WPI; 2004-305094/28.

Gierse JK;

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antidiabetic, immunomodulator, cardiant, neuroprotective, antinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention also describes: (1) antisense compounds having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                       /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 8 A; 8 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                  /*tag= c
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
                                                                                                                                              /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 1431; 132pp; English.
Location/Qualifiers
                                                     'mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                    25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                                                                                                                                                                        25-SEP-2002; 2002US-0413549P.
                                    Ω
                                                                                                                            *tag= a
                     . 50
                                                                                                                                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-305094/28
                                                                                                                                                                                                                                                                            WO2004028458-A2
Key
modified_base
                                                                                                       modified_base
                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                               08-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpCES-1). The human mpCES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpCES-1, which specifically Mybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpCES-1 in cells or tissues; and (3) a method of treating an animal mpCES-1 in cells or tissues; and (3) a method of treating an animal mpCES-1 in collisonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, cardiant, neuroprotective, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpCES-1 inhibitors and in gene therapy. The antisense compound to be used as mpCES-1 inhibitors and in gene therapy. The antisense compound condition associated with mPCES-1 e.g., inflammation, Alzheimer's chabetes, cancer, ischaemia or repertusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 1566; 132pp; English
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Gaps

; 0

3; Indels

0; Mismatches

Local Similarity 85.0

Best Loca Matches

2328 TGTGTGTGTGTGTGTGTG 2347

20 réreccérérérérerare 1

g

ADM15379 standard; DNA; 20 BP

RESULT 1399

01-JUL-2004 (first entry)

ADM15379;

ADM15379/ ID ADM1 XX AC ADM1 XX DT 01-J

Sequence 20 BP; 10 A; 8 C; 0 G; 2 T; 0 U; 0 Other;

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New compound of 8-80 nucleobases in length which inhibits the expression
                                                                                                                 se; fatty acid synthase; antisense therapy; metabolic rate; adiposity; serum leptin; serum cholesterol; blood glucose; serum insulin; serum lipid; breast cancer; prostate cancer; colon cancer; endometrium cancer; ovary cancer; thyroid cancer; infection; inflammation; tumour; probe; human.
                                                                                           Human fatty acid synthase, antisense oligonucleotide #45.
              ADO21340 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-OCT-2002; 2002US-00274085.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2002; 2002US-00274085.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-340035/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (FREI/) FREIER S M.
(DOBI/) DOBIE K W.
(BHAN/) BHANOT S.
                                                                                                                                                                                                                                                                                                                                                                                                                                     US2004077570-A1
                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                                             modified base
                                                                                                                                                                                                     Homo sapiens
                                                                 15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Freier SM,
                                       ADO21340;
 ADO21340/c
              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to antisense oligonucleotides targeted to a nucleic acid molecule encoding fatty acid synthase, which inhibit the expression of fatty acid synthase. The antisense oligonucleotides are useful for increasing the metabolic rate, decreasing adiposity, decreasing serum leptin, decreasing serum cholesterol, decreasing blood glucose, decreasing serum insulin, decreasing serum lipids and treating an animal having a disease or condition associated with fatty acid synthase, such as breast, prostate, colon, endometrium, ovary and thyroid cancers. It may also be useful prophylactically, e.g., to prevent or delay infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation or tumour formation. The present sequence represents a human fatty acid synthase antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New compound of 8-80 nucleobases in length which inhibits the expression
                                                                                                                                                                                                                                                        ss; fatty acid synthase; antisense therapy; metabolic rate; adiposity; serum leptin; serum cholesterol; blood glucose; serum insulin; serum cholesterol; blood glucose; serum insulin; eserum cancer; crostate cancer; colon cancer; endometrium cancer; ovary cancer; thyroid cancer; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of fatty acid synthase, useful for treating disease or condition associated with fatty acid synthase such as breast and colon cancers.
                            Gaps
                            ö
0.4%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 1.5e+03;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                               Human fatty acid synthase, antisense oligonucleotide #126.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 1 A; 8 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 16; SEQ ID NO 145; 87pp; English.
                                                                                                                                                                                                                                                                                                               inflammation; tumour; probe; human.
                                                     2326 rerererecererere 2345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bhanot S;
                                                                      |||||| || || ||||||||| TGTGTATGTGTGTG 1
                                                                                                                                                   ВБ
                                                                                                                                                                                                                                                                                                                                                                                                                        17-OCT-2002; 2002US-00274085
                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2002; 2002US-00274085
                                                                                                                                                  ADO21421 standard; DNA; 20
                                                                                                                                                                                                     (first entry)
                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Freier SM, Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-340035/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FREI/) FREIER S M.
            Local Similarity
les 17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DOBIE K W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BHAN/) BHANOT S.
                                                                                                                                                                                                                                                                                                                                                                   US2004077570-A1.
                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                    15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                             22-APR-2004.
                                                                              20
                                                                                                                                                                           AD021421;
 Query Match
                                                                                                                       RESULT 1400
               Best Loc
Matches
                                                                              요
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5

/note= "Phosphorothioate backbone. All cytidines are methylcytidines" $\ensuremath{\text{methylcytidines}}$

Location/Qualifiers

(first entry)

/*tag= b /mod_base= Other

/mod_base= Other /note= "2'-methoxyethyl (2'-MOB) nucleotides"

/note= "2'-methoxyethyl (2'-MOE) nucleotides" 16. .20

/mod_base= Other

/*tag=

æ *tag=

```
The invention relates to antisense oligonucleotides targeted to a nucleic acid molecule encoding fatty acid synthase, which inhibit the expression of fatty acid synthase. The antisense oligonucleotides are useful for increasing the metabolic rate, decreasing adiposity, decreasing serum leptin, decreasing serum cholesterol, decreasing blood glucose, decreasing serum insulin, decreasing serum lipids and treating an animal having a disease or condition associated with fatty acid synthase, such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     as breast, prostate, colon, endometrium, ovary and thyroid cancers. It may also be useful prophylactically, e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human fatty acid synthase antisense oligonucleotide.
of fatty acid synthase, useful for treating disease or condition associated with fatty acid synthase such as breast and colon cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 1.5e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 8 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                               Example 15; SEQ ID NO 64; 87pp; English.
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Bhanot

Dobie KW,

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Gaps

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1891 CTGCTGAAGGAGGCCACCG 1910

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Gaps

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0.4%; Score 15.2; DB 1; Length 20; 15.0%; Pred. No. 1.5e+03; ve 0; Mismatches 3; Indels

85.0%;

17; Conservative

Query Match Best Local Similarity

Best Loc Matches

GCCCTGAGTCTGGGCAGCTG 3647 GCCCTCAGTCTGGCTGCCG 20

3628

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RESULT 1401

15-JUL-2004 (first entry)

CTGCTGGAGCAGGGCCTCCG

20

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ss; fatty acid synthase; antisense therapy; metabolic rate; adiposity; serum leptin; serum cholesterol; blood glucose; serum insulin; serum lipid; breast cancer; prostate cancer; colon cancer; endometrium cancer; ovary cancer; thyroid cancer; infection;
                                                                                                                                                                                                                                                                                                                                                                                                          New compound of 8-80 nucleobases in length which inhibits the express of fatty acid synthase, useful for treating disease or condition associated with fatty acid synthase such as breast and colon cancers.
                                                                                             Human fatty acid synthase, antisense oligonucleotide #153.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fatty acid synthase antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 16; SEQ ID NO 172; 87pp; English
                                                                                                                                                                    tumour; probe; human.
                                                                                                                                                                                                                                                                                                                                                             Bhanot S;
                         ВР
                                                                                                                                                                                                                                                               17-OCT-2002; 2002US-00274085
                                                                                                                                                                                                                                                                                       17-OCT-2002; 2002US-00274085
                      ADO21448 standard; DNA; 20
                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                             Freier SM, Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-340035/31.
                                                                                                                                                                                                                                                                                                             (FREI/) FREIER S M. (DOBI/) DOBIE K W.
                                                                                                                                                                                                                                                                                                                                    (BHAN/) BHANOT S.
                                                                                                                                                                                                                 US2004077570-A1
                                                                                                                                                                  inflammation;
                                                                                                                                                                                           Homo sapiens
                                                                    15-JUL-2004
                                                                                                                                                                                                                                        22-APR-2004.
                                              AD021448
RESULT 1402
            AD021448
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expression

The invention relates to antisense oligonucleotides targeted to a nucleic acid molecule encoding fatty acid synthase, which inhibit the expression of fatty acid synthase. The antisense oligonucleotides are useful for increasing the metabolic rate, decreasing adiposity, decreasing serum leptin, decreasing serum cholesterol, decreasing blood glucose, decreasing serum insulin, decreasing serum lipids and treating an animal having a disease or condition associated with fatty acid synthase, such as breast, prostate, colon, endometrium, ovary and thyroid cancers. It may also be useful prophylactically, e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human

Gaps .. 0 ch 0.4%; Score 15.2; DB 1; Length 20; I Similarity 85.0%; Pred. No. 1.5e+03; 17; Conservative 0; Mismatches 3; Indels Query Match Best Local Similarity Matches 17; Conservat

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1891 CTGCTGAAGGAGGCCACCG 1910 crecregadcaeeeccrece 20

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ADO21309 standard; DNA; 20 BP AD021309 RESULT 1403 ADO21309/C ID ADO21309 XX AC ADO21309 XX

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New compound of 8-80 nucleobases in length which inhibits the expression
                                                                                                                                               /*tag= b
/mod_base= Other
/note= "Phosphorothioate backbone. All cytidines are 5-
methylcytidines"
                                          ss; fatty acid synthase; antisense therapy; metabolic rate; adiposity; serum leptin; serum cholesterol; blood glucose; serum insulin; serum lipid; breast cancer; prostate cancer; colon cancer; endometrium cancer; ovary cancer; thyroid cancer; infection; inflammation; tumour; probe; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of fatty acid synthase, useful for treating disease or condition associated with fatty acid synthase such as breast and colon cancers.
                                                                                                                                                                                                                  /mod_base= Other
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                    /*tag= c
/mod_base= Other
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                     Human fatty acid synthase, antisense oligonucleotide:#14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 33; 87pp; English;
                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                  Bhanot
                                                                                                                                                                                                                                                                                                                                                                17-OCT-2002; 2002US-00274085.
                                                                                                                                                                                                                                                                                                                                            17-OCT-2002; 2002US-00274085.
                                                                                                                                           1. .20
/*tag=
                                                                                                                                                                                                           /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                 Freier SM, Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-340035/31.
                                                                                                                                                                                                                                                                                                                                                                                    (FREI/) FREIER S M.
(DOBI/) DOBIE K W.
(BHAN/) BHANOT S.
                                                                                                                                                                                                                                                                                                JS2004077570-A1
                                                                                                                                         modified base
                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                           modified base
                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                      22-APR-2004
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ö The invention relates to antisense oligonuclectides targeted to a nucleic acid molecule encoding fatty acid synthase, which inhibit the expression of fatty acid synthase. The antisense oligonucleotides are useful for increasing the metabolic rate, decreasing adiposity, decreasing serum leptin, decreasing serum cholesterol, decreasing blood glucose, decreasing serum insulin, decreasing serum lipids and treating an animal having a disease or condition associated with fatty acid synthase, such as breast, prostate, colon, endometrium, ovary and thyroid cancers. It may also be useful prophylactically, e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human Sequence 20 BP; 4 A; 7 C; 8 G; 1 T; 0 U; 0 Other; fatty acid synthase antisense oligonucleotide. Local Similarity Query Match

Gaps ó ch 0.4%; Score 15.2; DB 1; Length 20; 1 Similarity 85.0%; Pred. No. 1.5e+03; 17; Conservative 0; Mismatches 3; Indels Best Loca Matches

ð a RESULT 1404

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hyperproliferative disorder (e.g. cancer), or conditions arising from bacterial or viral infections, or involving aberrant apoptosis. They are also useful in research and disgnostics for modulating the expression of cyclin-dependent kinase 6. The present sequence represents a cyclindependent kinase 6 antisense oligonucleotide. Note: Seqid 15-134 are also used in Tables 1 and 2 (page 30-34) but these sequences do not match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to antisense oligonucleotides targeted to cyclin-
dependent kinase 6, and which inhibit the expression of cyclin-dependent
kinase 6. The antisense oligonucleotides are useful for treating a
disease or condition associated with cyclin-dependent kinase 6, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding cyclin-dependent kinase 6, useful for treating cancer, bacterial/viral infection or conditions involving aberrant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                 Cyclin-dependent kinase 6, antisense oligonucleotide #114.
                                                                                                                                                 antisense therapy, cyclin-dependent kinase 6; hyperproliferative disorder; cancer; bacterial infection; viral infection; apoptosis; ss; probe; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 128; 68pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1610 AGTGCATCCACAGGGACCTG 1629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AGTICACCCACAGGCACCIG 1
              ADO32017 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                            31-JUL-2002; 2002US-00210802
                                                                                                                                                                                                                                                                                                                            31-JUL-2002; 2002US-00210802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   segid 15-134 of the seg list
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                   Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-356241/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
les 17; Conserv
                                                                                                                                                                                                                                                         US2004087523-A1.
                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                          06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                   Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         apoptosis
                                                ADO32017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1405
AD032017/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Loc
Matches
              8
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The invention relates to antisense oligonucleotides targeted to cyclindependent kinase 6, and which inhibit the expression of cyclin-dependent kinase 6. The antisense oligonucleotides are useful for treating a disease or condition associated with cyclin-dependent kinase 6, such as a hyperproliferative disorder (e.g. cancer), or conditions arising from bacterial or viral infections, or involving aberrant apoptosis. They are also useful in research and diagnostics for modulating the expression of cyclin-dependent kinase 6. The present sequence represents a cyclin-dependent kinase antisense oligonucleotide. Note: Seqid 15-134 are also used in Tables 1 and 2 (page 30-34) but these sequences do not match

Seguence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;

New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding cyclin-dependent kinase 6, useful for treating cancer, bacterial/viral infection or conditions involving aberrant

31-JUL-2002; 2002US-00210802 31-JUL-2002; 2002US-00210802

US2004087523-A1

06-MAY-2004

Homo sapiens.

(ISIS-) ISIS PHARM INC Freier SM, Dobie KW; WPI; 2004-356241/33. Disclosure; SEQ ID NO 74; 68pp; English.

apoptosis

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    S. pombe crm1 deletion cassette downstream tag BC775, SEQ ID NO:10.

                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lee MY;
                             ö
Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                Drug screening; fission yeast; Schizosaccharomyces pombe; deletion mutant; target gene; deletion cassette;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim HB,
                                                                                                                                                                                                                                                          deletion mutant; target gene; deletion cassette; homologous recombination; crml; tag sequence; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                               (KORE-) KOREA RES INST BIOSCIENCE & BIOTECHNOLOG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Huh GR, Jang YJ, Jung GS, Kim DU,
                                                     1610 AGTGCATCCACAGGGACCTG 1629
                                                                            1 AGTTCACCCACAGGCACCTG 20
                                                                                                                                            ADP70161 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                            13-DEC-2002; 2002KR-00079882
                                                                                                                                                                                                                                                                                                                                                                                                      03-MAY-2002; 2002KR-00024460
   Query Match 0.4%;
Best Local Similarity 85.0%;
Matches 17; Conservative
                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                          KR2003086216-A.
                                                                                                                                                                                               12-AUG-2004
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                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                     ADP70161;
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                                                                                                                    RESULT 1406
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                             such as a
                                                                                                                                                                                                Gaps
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antisense therapy, cyclin-dependent kinase 6; hyperproliferative disorder; cancer; bacterial infection; viral infection; apoptosis; ss; probe; human.

Cyclin-dependent kinase 6, antisense oligonucleotide #60

(first entry)

29-JUL-2004

24444444444

ADO31963

ВЪ

ADO31963 standard; DNA; 20

AD031963

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Gaps

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Synthetic.
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ADP76303
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                                                                                               The invention relates to a method for screening drugs using systematic deletion mutants of fission yeast (Schizosaccharomyces pombe). The deletion mutants exhibit a sensitive response to drugs and are created by deletion of a target gene selected from nda2, pspl, pakl, crml, ste<sup>7</sup>, stell, dis3, dis2, rad16, rad24, chkl, cdsl, cdrl and cdc25. Target gene deletion is achieved by homologous recombination with a target gene auxotrophic marker gene), tag sequences, binding sites for universal primers, a restriction site, and target gene specific sequences, primers, a restriction site, and target gene-specific sequences, primers, preferably those encoding N- and C-terminal sequences of the encoded protein. Sequences used in the
                                           Method for screening drugs using systematic deletion mutant of fission
                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                       Score 15.2; DB 1; Length 20; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                              3; Indels
                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 11 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                            construction of target gene deletion cassettes.
 Yoo HS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric phosphorothioate oligonucleotide #88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "2-methoxyethyl
                                                                           Example 2; SEQ ID NO 10; 28pp; Korean
Won MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                   875 ACGAGGCGGCCAGTGTAT 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= other
                                                                                                                                                                                                                                                                                                                                 20 ACGAGGGGGGGGGGGAGTGTAT 1
                                                                                                                                                                                                                                                                                                                                                                                               ADP76289 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                      0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-OCT-2002; 2002US-0419268P
ζĬ,
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                                                                                                                                                                                                                                                                                              17; Conservative
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 Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Crosby SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                   Local Similarity
                      WPI; 2004-199754/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-348453/32.
Park HO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reperfusion; ss.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Broschat KO,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                      ADP76289;
Lee YU,
                                                                                                                                                                                                                                                                                                                                                                          RESULT 1407
                                                        yeast.
                                                                                                                                                                                                                                                                                               Matches
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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MOE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
0
New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
6
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 1.5e+03;
tive 0; Mismatches 3; IndelB
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 2 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric phosphorothioate oligonucleotide #102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note = "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= other
/note= "2-methoxyethyl wing"
17. .20
/*tag= b
                                                                                                        Claim 4; SEQ ID NO 88; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3238 AGTTGGAGGTGATTCCAGTG 3257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 AGTIGGIGAIGATICCATIG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'mod_base= other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
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                                                                 ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP76303 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Broschat KO, Crosby
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            reperfusion; ss.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          apolipoprotein B; ApoB; cardiovascular; antiarteriosclerotic; antilipaemic; antidiabetic; anorectic; cardiant; vasotropic; hypotensive; anabolic; eating disorder; cytostatic; endocrine; vasotropic; neuroprotective; nootropic; lipid; cholesterol metabolism; hyporlipoproteinaemia; hyperlipidaemia; hyperlipidaemia; hyperlipidaemia; hyperlipidaemia; sexual ateliotic dwarfism; hyperthyroidism; hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             anorexia nervosa; Werner's syndrome; hepatoma; multiple myeloma; uraemia; impotence; obstructive liver disease; Alzheimer's; dementia; diabetes; obsesity; atherosclerosis; antiaense; 2'-MOB gapmer; 2'-methoxyethyl wing; phosphorothioate backbone; human; chromosome 2p21-2p24; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "OTHER = Phosphorothioate backbone, bases 1-5 and 16-20 2'-MOE wing bases, all cytidine residues are 5-methylcytidines"
(GFAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 4 A; 2 C; 5 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotides inhibit human GFAT expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                      Claim 4; SEQ ID NO 102; 175pp; English.
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15-MAY-2003; 2003WO-US015493.
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Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AD032937;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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AD032937
ID AD032937
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AD032937
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AD032937
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AD0329
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The invention relates to a novel antisense compound where the compound by the invention relates to an inhibits expression of mRNA encoding human applicable and inhibits expression of mRNA encoding human applicable in Culture at a concentration of 150 nM. The confluent HepG2 cells in culture at a concentration of 150 nM. The compound of the invention demonstrates cardiovascular, cardiant, vasotropic, antilipaemic, antidiabetic, anorectic, cardiant, vasotropic, neuroprotective and nootropic activities and may be useful for inhibiting the expression of apolipoprotein B in cells or tissues in vivo in order to address a condition associated with abnormal lipid or cholesterol metabolism. The compound may be useful for eactants and chylomicrons and thus may be utilised during treatment of hyperlipoproteinsemia, hyperlipidaemia, hypercholesterolaemia, cardiovascular disorders, Von Gierke's disease, lipodystrophy, Cushing's syndrome, sexual ateliotic dwarfism, hyperthyroidism, hypertension, anorexia nervosa, Werner's syndrome, hepatoma, multiple myeloma, uraemia, impotence, obstructive liver disease, Alzheimer's disease, dementia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                breast cancer; cytostatic; gene therapy; human; laminin alpha 4; LAWA4; chromosome 6q21; 88; PCR; primer; SNP; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence is that of
the invention which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                     Antisense oligonucleotide compound that inhibits expression of mRNA encoding human apolipoprotein B, useful for treating hyperlipidemia, diabetes, obesity, von Gierke's disease, lipodystrophies, Cushing's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer 1 used to genotype human laminin alpha 4 polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 15.2; DB 1; Length 20; 35.0%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    impotence, obstructive liver disease, Alzheimer's disease, diabetes, obesity and atherosclerosis. The current sequenc antisense 2'-MOE (2'-methoxyethyl) gapmer oligo of the inv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reneland R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Braun A, Kammerer SM,
                                                                                                                                                  Example 33; SEQ ID NO 385; 483pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2567 ACCACGGGACATCACAGGGT 2586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 ACCACCTGACATCACAGGTT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-NOV-2003; 2003WO-US037966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        82.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             targeted to human ApoB RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP08405 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 85.0 tes 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SEQU-) SEQUENOM INC.
                 WPI; 2004-420321/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-441082/41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roth RB,
                                                                                                              syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADP08405;
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Matches
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                          Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense therapy, endothelial lipase; endothelial lipase associated disorder; cardiovascular disease; mouse; antisense oligonucleotide; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
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/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
15. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                           Example 2; Page 73; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     374 GCATTGGAGGCATCAAGCTG 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hase= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 GCATTGTAGTCATCCAGCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP66864 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match
Best Local Similarity 85.0°
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-449390/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1411
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XX ADP6

XX ADP7

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XFFFXXXXCCCCCCCCXX
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The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding endothelial lipase. The compound compound specifically hybridises with the nucleic acid molecule encoding conducted living the such as a sequence of 3927 bp fully defined conducted are: inhibiting the expression of endothelial lipase. In the specification) and inhibiting the expression of endothelial lipase. In cells or tissues; screening for amodulator of endothelial lipase; a diagnostic method for identifying a disease state; a kit or assay device compution associated with endothelial lipase; compition associated with endothelial lipase, comprising administering to the animal a therapeutic or prophylactic amount of the compound so that conjugonucleotide is useful for inhibiting the expression of endothelial lipase is inhibited. The antisense of coligonucleotide is useful for inhibiting the expression of endothelial lipase expression, such as cardiovascular disease. In addition, the compound is used for diagnostics, prophylaxis, or as tresarch reagents or kits. This sequence represents a mouse endothelial lipase antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
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/mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense therapy, endothelial lipase, endothelial lipase associated disorder; cardiovascular disease; mouse; antisense oligonucleotide; antisense technology; ss.
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/mod_base= OTHER
/note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
             New antisense oligonucleotides for modulating endothelial lipase expression, useful for diagnosing, preventing or treating diseases associated with aberrant endothelial lipase expression, e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                          Example 16; SEQ ID NO 120; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               485 TCCGGCAGACGTACACGCTG 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 TCCTGCATACCTACACGCTG 1
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Matches 17; Conservative
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                                                                       cardiovascular disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1412
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ADP66991
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The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding endothelial lipase. The compound especifically hybridises with the nucleic acid molecule encoding endothelial lipase (which comprises a sequence of 3927 bp fully defined in the specification) and inhibits the expression of endothelial lipase. Also described are: inhibiting the expression of endothelial lipase; a clist or tissues; screening for a modulator of endothelial lipase; a diagnostic method for identifying a disease state; a kit or assay device comprising the above compound; and treating an animal having a disease or condition associated with endothelial lipase, comprising administering to the animal a therapeutic or prophylactic amount of the compound so that expression of endothelial lipase is inhibited. The antisense of endothelial lipase is inhibited. The antisense of endothelial lipase is inhibited. The antisense of endothelial lipase as consisted with a coligonucleotide is useful for inhibiting the expression of endothelial lipase expression, such as cardiovascular disease. In addition, the compound is used for diagnostics, prophylaxis, or as the endothelial lipase for diagnostics, prophylaxis, or as the endothelial lipase and the endothelial and the endothelial endothelial control or the endothelial endothelia
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                                                                                                                                                                                                                                                      New antisense oligonucleotides for modulating endothelial lipase expression, useful for diagnosing, preventing or treating diseases associated with aberrant endothelial lipase expression, e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                Example 16; SEQ ID NO 247; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  485 TCCGGCAGACGTACACGCTG 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   research reagents or kits. This s
lipase antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCCTGCATACCTACACGCTG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADQ09888 standard; DNA; 20 BP
12-DEC-2002; 2002US-00319915.
                                                  12-DEC-2002; 2002US-00319915.
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                                                                                                                                                                                                                                                                                                                                cardiovascular disease.
                                                                                                    (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                    WPI; 2004-449390/42.
                                                                                                                                                       Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AD009888;
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ADQ09888/c
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g
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polypopulue (analyof an isolateu nuclear actue depressing the prosperior) an insulin-producing cell comprising a vector (where the vector comprises a nucleic acid sequence encoding an islet cell differentiation teranscription factor), an insulin-producing cell (generated by a method comprising obtaining a somatic cell and transfecting the cell with a comprising a nucleic acid sequence encoding an islet cell with a confident transcription factor, where in the transfecting step the cell produces insulin), a method of generating at least one pancreatic cell produces insulin), amethod of generating at least one pancreatic coll produces insulin) (comprising: NeuroD or ngn3 polypeptide or a polymocleotide expressing a betacellulin polypeptide).

The islet cell differentiation transcription factor polypeptide is comprosed in a polymocleotide expressing a betacellulin polypeptide).

The islet cell differentiation transcription factor polypeptide is cortein), Nkx6.1, IB1-1 (Islet factor 1), Pdx-1 (pancreatic and duodenal conneobox protein), Nkx6.1, IB1-1 (Islet factor 1), Pdx-1 (pancreatic and duodenal cell differentiation transcription factor oplypeptide or its homologue canalogue is useful for treating a mammal for insulin-dependent diabetes, contracting an insulin manner of the paragraphy of paragraphy

The invention relates to the use of an islet cell differentiation transcription factor polypeptide or its homologue or analogue for treating a mammal for insulin-dependent diabetes, increasing an insulin level in a somatic cell, or generating an insulin-producing cell. Also included are a method of treating a mammal for insulin-dependent diabetes, a method of increasing an insulin level in a somatic cell, a method of generating an insulin level in a somatic composition comprising an isolated isset cell differentiation transcription factor polypeptide (and/or an isolated nucleic acid expressing the polypeptide).

Use of an islet cell differentiation transcription factor polypeptide or tits homologue or analog for treating a mammal for insulin-dependent diabetes, increasing an insulin level in a somatic cell, or generating an insulin-producing cell.

(BAYU) BAYLOR COLLEGE MEDICINE.

WPI; 2004-517032/49. Chan L, Kojima H;

03-SEP-2002; 2002US-0407743P.

Example 20; SEQ ID NO 126; 190pp; English

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                                                                                                                                                                                                                                                                                                                                                                   producing cell. The present sequence is a reverse transcriptase PCR (RT-PCR) primer used to study the expression of an islet cell differentiation franscription factor polynucleotide for the proteins detailed above in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue-specific expression; epithelium; carcinoma; metabolic; disorders; phenylketonuria; cystic fibrosis; haemophilia; emphysema; vaccine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.5e+03; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 5 C; 7 G; 6 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1609 AAGTGCATCCACAGGGACCT 1628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 AAGCGCATCCACAGGGCCAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ66994 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 85.0 les 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DF3 enhancer fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
18-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ66994;
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1414
                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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insulin-dependent diabetes; insulin; somatic cell; insulin-producing cell; islet cell; NeuroD; ngn3; neurogenin 3; Pax6; paired-box transcription factor 6; Pax4; Pax4; paired-box transcription factor 4; Nkx2.2; NK-2 class homeobox protein; Nkx6.1; Isl-1; Islet factor 1; Pdx-1; Pax1; Pax1; Pax1; R-PCR; reverse transcription copy protein; BCT; betacellulin; primer; RT-PCR; reverse transcriptes PCR.

03-SEP-2003; 2003US-00654102.

US2004132679-A1.

08-JUL-2004

Mus musculus

Mouse; 88; PCR; islet cell differentiation transcription factor;

Mouse RT-PCR primer set Insulin-2, forward primer.

(first entry)

23-SEP-2004

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Gaps

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primer (AAQ73753) and a 3'-primer (AAQ73754) corresponding to nucleotides 4-23 and 995-1115, respectively, of the promoter sequence. The promoter can be operatively linked to the branching enzyme gene or to heterologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Specifically designed oligodeoxyribonucleotides form triplexes in single-
or double-strand DNA at homopurine-homopyrimidine targets. These
triplexes block in vitro NNA synthesis by all DNA polymerases studied,
including Sequenase3, Taq, Vent, and Pol I. A similar phenomenon occurs
when DNA polymerases are supplemented with accessory replication
proteins, including SSB protein. Replication blockage is highly sequence-
specific and even one or two point substitutions within either the target
sequence or the oligonucleotide abolish the effect. Sequence-specific
blocking of DNA replication in vivo is facilitated by the methods and
compositions of the present invention. The present sequence is a triplex-
forming oligonucleotide which targets a 185 bp long stretch within the
inverted repeat (IR-L) of herpes simplex virus 1 at position 117156-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence specific inhibition of DNA synthesis - by triplex-forming oligo:nucleotide(s), for detection of oncogene mutation(s) and treatment of e.g. HSV, Hepatitis C and Papillomavirus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    HBV; oligodeoxyribonucleotide; homopurine-homopyrimidine target; block; in vitro; DNA synthesis; DNA polymerase; Sequenase3; Taq; Vent; Pol I; accessory replication protein; SSB protein; sequence-specific; triplex forming oligonucleotide; exon 3; inverted repeat; IR110; hepatitis B virus; P gene; ss.
                                                                                                                     Query Match 0.4%; Score 15.2; DB 1; Length 21; Best Local Similarity 85.0%; Pred. No. 1.6e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 0.4%; Score 15.2; DB 1; Length 21; 1 Similarity 85.0%; Pred. No. 1.6e+03; 17; Conservative 0; Mismatches 3; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                        Triplex-forming oligonucleotide targetting HSV-1 IR-L.
                                                                                      Sequence 21 BP; 0 A; 3 C; 12 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 21 BP; 0 A; 0 C; 15 G; 6 T; 0 U; 0 Other;
                                                        genes for expression in plant seeds
                                                                                                                                                                                             2319 GTGTGTGTGTGTGTGTGT 2338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 40; 78pp; English.
                                                                                                                                                                                                                            érérérégérérécécéde 21
                                                                                                                                                                                                                                                                                                                     BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UNII ) UNIV ILLINOIS FOUND
                                                                                                                                                                                                                                                                                                                   AAT35039 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-300649/30.
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Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9618732-A2
                                                                                                                                                                                                                                                                                                                                                                                      18-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mirkin SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                    AAT35039;
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                                                                                                                                                                                                                                                                                                                                                                                    can be linked to a heterologous coding sequence to increase tissue-
specific expression of the coding sequence in the spitchilum. These
prods. can be used to treat carcinomas or metabolic disorders such as
phenylketonuria, cystic fibrosis, haemophilia or emphysema. They can also
be used as vaccines against infectious organisms or can confer resistance
to the effects of a chemotherapeutic drug. See also AAQ66988-95. (Updated
on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                               Isolated DF3 enhancer DNA sequence - is used to enhance epithelium tissue -specific expression for treating carcinomas or metabolic disorders.
                                                                                                                                                                                                                                                                                                                                                                    The sequence represents a DF3 gene 5' flanking fragment. The DF3 enhancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The rice starch branching enzyme gene promoter was amplified using a 5'-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Starch branching enzyme promoter; rice; starch content; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New gene of branching enzyme of rice starch - useful for increasing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 4 A; 0 C; 15 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rice starch branching enzyme promoter 3'-primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MITS-) MITSUI GYOSAI SHOKUBUTSU BIO KENKYUSHO.
                                                                                                                                                                          (DAND ) DANA FARBER CANCER INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3067 TCCCACACCCCAACACTTCC 3086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 13; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 29; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              recerececeaceaerree 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ73754 standard; DNA; 21 BP
                                                                                                        93WO-US011915
                                                                                                                                           92US-00999742
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17; Conservative
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                                                                                                                                                                                                                                              WPI; 1994-249205/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                              Abe M;
                                    WO9416057-A1
                                                                                                       08-DEC-1993;
                                                                                                                                           31-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP06261767-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JUL-1995
                                                                      21-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-SEP-1994
   Synthetic.
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                                                                                                                                                                                                              Kufe D,
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the expression of a target nucleic acid (tNA) sequence via binding of the compounds with the tNA sequence. The method comparises generating a library of virtual compounds in silico according to defined criteria, and evaluating in silico the binding of the virtual compounds with the tNA according to defined criteria. Also described are: (1) a method of defined criteria, and evaluating in silico the binding of the virtual compounds with the tNA according to defined criteria. Also described are: (1) a method of a tNA sequence via binding of the ONS with the tNA sequence comprising generating a library of virtual compounds in silico according to defined criteria; and evaluating in silico the binding of the virtual ONS with the tNA according to defined criteria; and (2) a method of defining a set of compounds that medulate the expression of a tNA sequence via binding of the compounds with the tNA. The methods can be used for the generation and identification of synthetic compounds having defined physical, chemical or bloactive properties. Information gathered from assays of tractable to a variety of nucleotide sequences that are tractable to a variety of nucleotide sequence based technologies, e.g. antisense drug discovery and target validation. AAZ40852 to AAZ41220, and ANZ20011 to AAX52106, represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying compounds which modulate expression of nucleic acids, used to provide compounds having defined physical, chemical or bioactive properties, e.g. antisense activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sasmor HM, Brooks DG;
                                                                                                                                                                                                                                                                                             Identification; genetic target; gene modulation; human; probe; antisense oligonucleotide; phosphorothioate; PCR primer; nucleotide sequence-based technology; antisense drug discovery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 3 A; 10 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Freier SM,
Vickers TA;
                                                                                                                                                                                                                                                          Human ELK-1 PCR reverse primer SEQ ID NO:203.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 23; Page 103; 264pp; English.
2922 GCGGGCGTGGGGGGCGTG 2941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ?, Mcneil J,
Borchers AH,
                                   2 crecercrerececerere 21
                                                                                                                                      AAZ41051 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US008268.
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98US-00067638
                                                                                                                                                                                                                   26-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cowsert LM, Baker BF,
                                                                                                                                                                                                                                                                                                                                                         target validation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ohasi C, Wyatt JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-620446/53.
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9953101-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-APR-1998;
28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                             AAZ41051;
                                                                                               RESULT 1417
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98WO-US019419

17-SEP-1998;

17-SEP-1997; 09-JUN-1998;

98US-00093972 97US-0059160P

EAST CAROLINA.

(UYEC-) UNIV

Nyce JW;

WPI; 1999-229400/19.

Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vascoonstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impaded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vascoonstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;

prostate cancer; ss.

WO9913886-A1.

Synthetic.

C/EBP-beta antisense oligonucleotide fragment

05-JUL-1999 (first entry)

AAX55050;

AAX55050 standard; DNA; 21 BP.

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The specification describes antisense oligonucleotides (AAX52869-X55271)

directed against at least 2 mRNAs selected from target genes, coding and
non-coding regions of RNAs corresponding to target genes, coding and
codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'
end and the juxta-section between coding and non-coding regions and all
segments of RNAs encoding proteins associated with one or more diseases,
conditions or mixtures. The antisense coligonucleotides may be derived
from sequences AAX5512-4. These multiple target oligonucleotides
from sequences AAX55180-271) can be used for the antisense treatment of
diseases and conditions. Typical diseases and conditions are those
associated with impaired respiration and inflammation, including lung
diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis,
cute asthma, allergies, asthma, impeded respiration, respiratory
distress syndrome, pain, cystic fibrosis, pulmonary hypertension,
culmonary vasoconstriction, emphysema, chronic obstructive pulmonary
distress (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g.
colon cancer, breast cancer, lung cancer, parcer, cancer, papatic metastases,
hepatocellular carcinoma, kidney cancer, metastasize or have metastasized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the lungs, including breast and prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 0 A; 6 C; 14 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 70; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03; tive 0; Mismatches 3; Indels

Best Local Similarity 85.0 Matches 17, Conservative

Query Match

New antisense oligonucleotides used in treatment of, e.g. pulmonary vasoconstriction.

(first entry)

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Lymphocytic virus-binding lectin LVBL - and corresponding DNA, vectors, antibodies, etc., useful for diagnosis or therapy of HIV infection.
                                                                                                                                                                                                                                                         Human, lymphocytic virus-binding lectin, LVBL, HIV, gene expression,
antibody; primer, probe, diagnosis, infection, vector, antisense,
cell proliferation, haematopoietic; progenitor cell, aplasia, 88;
                                                                                                                                                                                                            Human Lymphocytic Virus Binding Lectin gene primer #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYNI-) UNIV NICE-SOPHIA ANTIPOLIS.
                                                       AAX79319 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                      medullary hypoplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-349542/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lefebvre JC,
                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-DEC-1997;
                                                                                                                                                         31-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                          AAX79319;
  RESULT 1420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a composition comprising two oligonucleotide analogues, each having a binding domain and a coupling moletty, where the binding domains are capable of hybridizing to a target polynucleotide and the coupling moleties are capable of coupling to each other in the absence of a target molecule. The composition/compound is used to cleave an RNA target. The compositions can be used to determine an optimal antisense site for a given mRNA or an optimal ribozyme cleavage site for a target RNA. By separating the antisense molecules into two or more pieces, a comprehensive antisense library can be prepared in advance, rather than synthesizing a plurality of candidate antisense molecules as needed. A complete library of every possible 17-mer oligonucleotide, using the four natural bases, would consist of 417 or 1010 molecules. By providing the antisense molecules in at least two components, e.g. a library of 8-mers and a library of 9-mers, assembled quickly as needed, the library size is reduced to 48 + 49, or 327 650 molecules. The complexity of the library can be further some of the natural bases. The present sequence represents an oligonucleotide, which is used in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide analog compositions capable of coupling to form antisense
       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                     Conbinatorial antisense library; oligonucleotide analogue; RNase; ribozyme; cleavage; anchor; binding; target RNA; ss.
       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.2; DB 1; Length 21; 70.0%; Pred. No. 1.6e+03; ive 3; Mismatches 3; Indels
     Indels
                                                                                                                                                                                                                                                                                                                                                                                     BCL2-targeted antisense oligonucleotide SEQ ID NO:49.
       3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 3 A; 6 C; 6 G; 1 T; 5 U; 0 Other;
0; Mismatches
                                                     1476 GCGGCCCGCGCCCCCCG 1495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 9; Page 46; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arnold LJ;
                                                                                       6066666666666666666
                                                                                                                                                                                                                                 AAX33518 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-0060673P.
98US-00136080.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-US020361
                                                                                                                                                                                                                                                                                                                                   (first entry)
  17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brown BD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-264039/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (OASI-) OASIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9918238-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-AUG-1998;
                                                                                                                                                                                                                                                                                                                                   07-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Riley TA,
                                                                                                                                                                                                                                                                                   AAX33518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                               RESULT 1419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loca
Matches
  Matches
                                                                                                                                                                                                      AAX33518

AAX33518

AAX33518

AAX3518

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Giordanengo V, Bannwarth

97FR-00015224.

97FR-00015224

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                                                       Sequences AAX79317-X79320 represent primers useful for detecting the gene
                                                                             encoding a human lymphocytic virus-binding lectin (LVBL, AAX79316). The encoding a human lymphocytic virus-binding lectin (LVBL, AAX79316). The lectin was isolated from HIV-infected CEM cells and has a molecular weight of 47 kD. LVBL gene expression was shown to be enhanced in the presence of an infective HIV strain. Thus antibodies to LVBL can be used to detect the LVBL polympotide and LVBL polymucleotides can be used as primers and probes for detecting LVBL molleic acids, e.g. for diagnosis of a viral infection, especially an HIV infection. Antibodies, vectors and (anti) sense oligonucleotides can be used to prevent or treat a viral infection, especially an HIV infection, or modulate cell proliferation, especially to stimulate proliferation of haematopoietic progenitor cells in the treatment of medullary hypoplasia or aplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human ELK-1; p62TCF; Ets domain transcription factor protein; apoptosis; expression inhibition; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.2; DB 1; Length
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reverse PCR primer for amplification of human ELK-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 8 A; 0 C; 13 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     184 GGGGAGGACGAGGA 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGGGAGGAAGAGGAGGA 20
Claim 12; Page 56; 72pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ06610 standard, DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 85.0°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ06610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ06610/
ID AAZ0
XX
AC AAZ0
XX
DT 23-N
XX
KW Huma
KW Huma
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; 0

Gaps

. 0

1434 GCTGGTGGAGTACGCGGCCA 1453

Conservative

1 GCUGGUUGAGUACUCCGCCA 20

WO200009525-A2.

99WO-US017712. 98US-0095212P

03-AUG-1999; 33-AUG-1998;

24-FEB-2000

(UYEC-) UNIV EAST CAROLINA

WPI; 2000-205971/18.

Nyce JW;

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AZ06608. Human ELK-1 also known as p62TCF is a member of the terrary complex factor subfamily of Ets-domain transcription factor proteins. Ar106081-206607 inhibit the expression of ELK-1 welleicacid molecule AAZ06571-206607 all cause at least 30% inhibition of ELK-1 expression. The antense sequences can be used to inhibit the expression of human ELK-1 in human cells or tissues in vitro. ELK-1 uses a bipartite recognition mechanism mediated by both protein-DNA and protein-protein interactions to control various signal transduction pathways and other cell functions including apoptosis. This means that antisense compounds inhibiting expression of ELK-1 can be used to treat disease associated with its expression in animals, particularly humans and to prevent or delay infection, inflammation or tumour formation. The compounds can also be used for diagnosis, as research reagents and in kits
                                                                                                                                                                                                                                                                                                                                                                                                                                                   primers AAZ06609-Z06610 are used to amplify the human ELK-1 sequence
                                                                                                                                                                                                                                                                                                                                                    Antisense compound useful for diagnosis, treatment and prevention of disease associated with ELK-1 expression.
diagnosis; phosphorothioate; antisense compound; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 3 A; 10 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                              Example 13; Col 37; 31pp; English
                                                                                                                                                                 98US-00213767.
                                                                                                                                                                                                      98US-00213767.
                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                Cowsert LM;
                                                                                                                                                                                                                                                                                                                  WPI; 1999-517959/43.
                                                    Homo sapiens
                                                                                                                                                                 17-DEC-1998;
                                                                                                                                                                                                      17-DEC-1998;
                                                                                         US5948680-A
                                                                                                                              07-SEP-1999
                                                                                                                                                                                                                                                                                Baker BF,
                                  Synthetic
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New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hyperrension, bronchitis, emphysema, respiratory distress syndrome, ischemia or

Disclosure; Page 539; 1343pp; English.

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The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets conclete acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antivillammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating ceffects afflict the lungs of a subject. They can be used for treating impaired respiration, respiratory desconsitiction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cysic fibrosis, pulmonary hypertension, emphysema, chronic obstructive continuously hypertension, emphysema, chronic obstructive carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the Non reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing conchoconstriction and inflammation. AAA3313 to AAA35312 represent the invention, which correspond to SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:1 to 185, but the sequences creations, named sequences sEQ ID NO:1 to 185, but the present invention. N.B. Sequences given in the disclosure of the present invention. N.B. Sequences given in the disclosure of the present invention. N.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 0 A; 6 C; 14 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human C/EBP polynucleotide fragment #2186.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1476 GCGGCGCGCCCCCGG 1495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 GCGGGGCGCGCGCCTGG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF20619 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF20619;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ð
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Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorochioate; impaired respiration; inflammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.

Homo sapiens

Human adenosine receptor related polynucleotide SEQ ID NO:2186.

(first entry)

28-JUL-2000

AAA34497;

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3; Indels

0; Mismatches

82.08;

Local Similarity 85.0

Matches

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Query Match

860 AGCTGGTGGAGGCTGACGAG 879

20 AGCTGGTGGATGCAGAGGAG 1

AAA34497 standard; DNA; 21 BP

RESULT 1422

0.4%; Score 15.2; DB 1; Length 21; B5.0%; Pred. No. 1.6e+03;

02-FEB-2001

AAC73548;

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The present invention describes low adenosine (A) content antisense oligonuclectides and compositions (I) comprising them. In the antisense coligonuclectides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonuclectides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with ungivespiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoslobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and chemokine receptors, adenosine receptors, bradykinin receptors, central creceptors, adenosine receptors, bradykinin receptors, central creceptors, adenosine receptors, bradykinin receptors, central creceptors, defensins, growth factors, vascative peptides and receptors, bradykinin receptors central creceptors, binding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or lung inflammation, allergic syndrome condition selected from pulmonary vasoconstriction, inflammation, cultanged respiratory vasoconstriction, inflammation, condition, cystic fibrosis (CP), allergic rhinitis (AR), pain, cystic fibrosis (CP), allergic rhinitis (AR), pain, cystic fibrosis (CP), cystic fibrosis (CP), cystic fibrosi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.
                              respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
  surfactant hypoproduction; pulmonary vasoconstriction, asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 14; Page 264; 1592pp; English.
                                                                                                                                                                                                                                                                                                                      24-MAR-2000; 2000WO-US008020.
                                                                                                                                                                                                                                                                                                                                                                          99US-0127958P
                                                                                                                                                                                                                                                                                                                                                                                                                        (UYEC-) UNIV BAST CAROLINA.
(NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-679539/66.
                                                                                                                                                                                                                   WO200062736-A2.
                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                          06-APR-1999;
                                                                                                                                                                                                                                                                    26-OCT-2000.
                                                                                                                cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW;
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pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                                                                                                                                                       0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                        Sequence 21 BP; 0 A; 6 C; 14 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                               Query Match
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1476 GCGGGCGCGCCCCCGG 1495
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17; Conservative

Best Local Similarity Matches 17; Conser

AAC73548 standard; DNA; 21 BP. RESULT 1424 AAC73548 ID AAC73548

NOV; olfactory; cytostatic; immunomodulator; vulnerary; anti-HIV; antiasthmatic; antiinflammatory; gastrointestinal; neuroprotective; osteopathic; gene therapy; odorant receptor; olfactory receptor; of-protein coupled receptor; ofCR; neuro-olfactory; trauma; probe; neoplastic disorder; cancer; adenocarcinoma; lymphoma; prostate cancer;

BP.

AAH75783 standard; DNA; 21

(first entry)

15-OCT-2001

AAH75783;

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Gaps

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Human NOV 12 probe.

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The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one such polymorphic locus used in the present invention. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to form the oligonucleotide array. Note: This sequence includes a SNP represented by the degenerate codon in the
                                                                                                                    Oligonucleotide array; genotyping; single base extension reaction; SBE; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping
                                                                                                                                                                                                                                                                                                                                                                                                         Lander ES, Lockhart DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .;
0
                                                                                   SNP flanking sequence #120 used in multiplexing PCR/SBE assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 7 A; 4 C; 5 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         Hirschhorn JN, Huang X, Kaplan P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            using single base extension reactions.
                                                                                                                                                                                                                                                                                                                                                    (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; Page 61; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    754 CACAACGICACCITIGAGGA 773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CACAAGGTCARCATTGAGGA 20
                                                                                                                                                                                                                                                                     27-MAR-2000; 2000WO-US008069.
                                                                                                                                                                                                                                                                                                      99US-0126473P.
                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 85.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                                                                                                                                                         Sklar P;
                                                                                                                                                                                                     WO200058516-A2.
                                                                                                                                                                      Unidentified.
                                                                                                                                                                                                                                                                                                      26-MAR-1999;
                                                                                                                                                                                                                                                                                                                        23-JUN-1999;
                                                                                                                                                                                                                                     05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                       Ryder T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                         Fan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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/*tag= a /standard_name= "single nucleotide polymorphism"

Lander ES;

99US-0167334P

Location/Qualifiers

vivlemore401-10.rng

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New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics,
                                          Single nucleotide polymorphism; SNP; human; cancer; inflammation; heart disease; paternity testing; forensic science; ds.
             NFE2L1 polymorphism containing DNA fragment #297.
                                                                                                                                                                                                                                                                                                                                                                                                                        including polymorphic sites, for phenotype correla
paternity testing, medicine and genetic analysis.
                                                                                                                                                                                                                                                                                                                     (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 53; 80pp; English.
                                                                                                                                                                                                                                                            17-NOV-2000; 2000WO-US031639
                                                                                                                                                                                                                                                                                                                                                Cargill M, Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-367705/38.
                                                                                                                                                                                                    WO200138576-A2
                                                                                                                                                                                                                                                                                        24-NOV-1999;
                                                                                                   Unidentified
                                                                                     sapiens
                                                                                                                                                                                                                                 31-MAY-2001.
                                                                                                                                             variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1427
                                                                                    Ношо
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to novel human NOVX proteins and coding sequences, where X is any number from 1 to 18 (see AAH75716-AAH7573), and AAG64400 and AAG66322-AAG66338). NOVX are members of the AAH75733, and codorant/Olfactory receptor (ONY family, which are G-protein coupled receptors (GPCRs). The NOVX proteins and coding sequences are useful as therapeutics, particularly in the manufacture of a medicament for treating a syndrome associated with a human diseases/disorders of the neuro-olfactory system, e.g. those induced by trauma, surgery and/or neoplastic disorders. Purthermore, the coding sequences and proteins are useful in treating cancer e.g. adenocarcinoma, lymphoma, prostate cancer, uterus cancer, inappropriate immune response, AIDS, asthma, Crohn's disease, multiple sclerosis or AAbright hereditary oteodystrophy. The coding sequences are also useful in gene therapy for treating the above conditions. The present probe was used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                               New NOVX polypeptides and polynucleotides, useful for treating or preventing a syndrome associated with a human disease (e.g. disorders of the neuro-olfactory system), as well as in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                           Vernet CAM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
 uterus cancer; immune response; AIDS; asthma; Crohn's disease; multiple sclerosis; Albright hereditary osteodystrophy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
iive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                          Li L,
                                                                                                                                                                                                                                                                                                                                                                                           Spytek KA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 229; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1182 GCCCGGCTGACCCTGGGCA 1201
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31-JUL-2000; 2000US-0221943P.
21-DEC-2000; 2000US-0257599P.
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27-JAN-2000; 2000US-0178409P.
27-JAN-2000; 2000US-0178413P.
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07-FEB-2000; 2000US-0180634P.
24-JUL-2000; 2000US-0220516P.
                                                                                                                            29-JAN-2001; 2001WO-US002849
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Macdougall J;
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                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-514556/56.
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Best Local Similarity
                                                                      WO200155179-A2
                                                                                                                                                                                       27-JAN-2000;
                                                                                                                                                                                                                                                                                                                                    38-JAN-2001;
                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                          SK,
                                                                                                 02-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                         Peyman JA,
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DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs) A method is included in the invention for analysing a nucleid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection heart microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as
                                                                                                                                                                                                                                                  a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
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                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 5 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 15.2; DB 1;
85.0%; Pred. No. 1.6e+03;
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(revised)
(first entry)

09-SEP-2004 12-SEP-2001

AAH62396; RESULT 1426
AAH62396
ID AAH62396
XX
AC AAH62396
AC AAH62396
DT 09-SEP-2
DT 12-SEP-2

AAH62396 standard; DNA;

Key

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DNA sequences AAH62100 - AAH6268B represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites of diagnosis or monitoring any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaccutical. SNP containing oligonucleotides are useful in medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
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/*tag= a
/standard_name= "single nucleotide polymorphism"
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/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                               Lander ES;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 73; 80pp; English
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Best Local Similarity 85.0
Matches 17; Conservative
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Unidentified.
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12-SEP-2001
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  variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analyzing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites of the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in embalications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
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/standard_name= "single nucleotide polymorphism"
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85.0%; Pred. No. 1.6e+03;
tive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                       (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                     Cargill M, Ireland JS, Lander ES;
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(first entry)
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                                                                                                                                                                                              WO200138576-A2.
                  Homo sapiens
Unidentified
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Unidentified,
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12-SEP-2001
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Query Match

Best Loc Matches

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RESULT 1428

AAH62649

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DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neonate screening; prenatal screening; gene chip; diagnosis; phenylketonuria; maple syrup disease; galactosemia; homocysteinuria; mediuu-chain acyl-coA-dehydrogenase deficiency; biotinidase deficiency; familial hypercholesterolemia; familial defective apolipoprotein-B; cystic fibrosis; Marfan syndrome; Smith-Lemil-Opitz syndrome;
                                                                                                                                                                                                                          New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
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Pred. No. 1.6e+03;
0; Mismatches 3; Indels
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Local Similarity 85.0%; Pred. No. 1.6e
hes 17; Conservative 0; Mismatches
                                                                                                                   (WHED ) WHITEHEAD INST BIOMEDICAL RES.
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                                                                                                                                                                                                                                                                                                  Claim 1; Page 68; 80pp; English.
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                                                 17-NOV-2000; 2000WO-US031639
                                                                                    99US-0167334P
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                31-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Virus with specific reproduction in a tumor well and effective expression of tumor angiogenesis inhibitor and its construction method.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human tumour vascular genesis inhibiting factor related PCR primer 31
                                                                                                    DNA chip, useful for neonatal or prenatal screening for many genetic diseases simultaneously, carries oligonucleotides complementary to phenotypically relevant reference sequences.
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85.0%; Pred. No. 1.6e+03;
tive 0; Mismatches 3; Indels
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                                                                                                                                                                            Claim 4; Page 71; 101pp; German.
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Matches 17; Conservative
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                                                                   WPI; 2001-457616/49.
(SEED/) SEEDORF U.
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                                  Cullen P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH79135;
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Gaps

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Disclosure; Page 34 (Disclosure); 52pp; Chinese

09-JAN-2001; 2001WO-EP000139 21-JAN-2000; 2000DE-01002446

(CULL/) CULLEN P.

WPI; 2001-596837/67.

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vivlemore401-10.rng

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The invention relates to developing a virus with highly-effective expression of tumour vascular genesis inhibiting factor. The virus comprises a nucleotide sequence encoding the vascular genesis inhibiting factor inserted into a non-essential proliferation region of a virus gene group. The virus selectively proliferates in a tumour cell such that along with replication of virus, the nucleotide sequence copy number of the encoded vascular genesis inhibiting factor is increased. The expression of the vascular genesis inhibiting factor inhibits tumour vascular development and inhibits formation, growth and transfer of tumour. The present sequence is that of a PCR primer, useful to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; FCTRX; myelogenous leukaemia; carcinoma; melanoma; glioma; ss; astrocytoma; congenital neonatal alloimmune thrombocytopaenia; infection; neurological disorder; neurodegenerative disorders; nerve trauma; familial myelodysplastic syndrome; Charcot-Marie-Tooth neuropathy; demyelinating Gardner syndrome; familial myelodysplastic syndrome; mental health condition; immunological disorder; allergy; infertility; bronchial asthma; Avellino type eosinophilia; lung disease; deafness; reproductive disorder; sprocherin Ia deficiency; desmoid disease; turcot syndrome; liver cirrhosis; hepatitis C; virucide; gastric disorders; pancreatic disease; Schistosoma mansoni infection; Spinocerebellar ataxia; Plasmodium falciparum parasitaemia; diabetes; corneal dystrophy-Greonow type I; Corneal dystrophy-lattice type I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reis-Bucklers corneal dystrophy; cytostatic; immunosuppressive; antiallergic; antiasthmatic; antiinflammatory; probe; antidiabetic; protozoacide; hepatorropic; virucide; ophthalmological; gynaecological; antiinfertility; immunostimulant; auditory; haemostatic; gene therapy; FCTRS; human complement CIR component precursor.
                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                       0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                 Sequence 21 BP; 9 A; 6 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                     1457 GTAACCTGCGGGAGTTTCTG 1476
                                                                                                                                                                                                                                                                                                                                                                                                                21 GTTACCTGCTGGATTTTCTG 2
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03-MAR-2000; 2000US-018671BP.
06-MAR-2000; 2000US-0187233P.
06-MAR-2000; 2000US-0187294P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-MAR-2000; 2000US-0190400P.
07-APR-2000; 2000US-0196018P.
03-JAN-2001; 2001US-0259548P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human FCTR5 DNA probe Ag427.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS14103 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                          Local Similarity 85.0
les 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
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Macdougall J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2001
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                                                                                                                                                                                                                           invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS14103;
                                                                                                                                                                                                                                                                                                         Query Match
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Mishra V, Mezes PS,

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carcinomas, melanomas, astrocytomas, congenital neonateal alloimmune thrombocytopaenia, neurological disorders, neurodegenerative disorders, neurodegenerative disorders, neurodegenerative disorders, neurodegenerative rooth neuropathy, demyelinating dardner syndrome, familial myelodysplastic syndrome, mental health conditions, immunological disorders, allergy and infection, bronchial asthma, Avellino type cosinophila, lung diseases, reproductive disorders, infertility, male and female reproductive disorders, desfiness, glycoprotein la deficiency, desmoid disease, turcot eyndrome, liver cirrhosis, hepatitis C, gastric disorders, pancreatic diseases such as diabetes, Schistosoma mansoni infection Spinocerebellar ataxia, Plasmodium falciparum parasitaemia, formeal dystrophy-Grenonuw type I, Corneal dystrophy-Grenonuw type I and Reis-Bucklers corneal dystrophy-This sequence represents a probe in DNA encoding FCTRS, a protein bearing sequence homology to human complement
                                                                                                                                           The invention relates to human FCTRX polypeptides, FCTR1-FCTR7, and the nucleic acids encoding them. These sequences are useful for the treatment or prevention of numerous disorders including myelogenous leukaemia,
                                        Novel polypeptides designated as FCTRX polypeptides, useful in detection, prevention and treatment of a broad range of pathological states.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hormone-regulated cancer, antisense oligonucleotide, IGFBP-2; insulin-like growth factor binding protein-2; hormone-regulated tumour; breast cancer; prostate cancer; IGF-1-sensitive cancer; apoptosis; hormone-responsive cancer; hormonal withdrawal; oligodeoxynucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Composition for treating hormone-regulated cancer, particularly of prostate or breast, comprises oligonucleotide antisense to insulin-like growth factor binding protein-2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic antisense IGFBP-2-oligodeoxynucleotide (ODN) #44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 3 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rennie PS;
                                                                                                    Example 2; Page 198; 215pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1182 GGCCCGGCTGACCCTGGGCA 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ODN; endocrine tumour therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 GGCCCAGCTGACCCTGCTCA 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK70356 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-2001; 2001WO-US028748.
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Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C1R component precursor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-339861/37.
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ABX15381;
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                 The present invention relates to a new composition for treating hormonerequiated cancer. The composition comprises an antiense oligonuclectide that inhibits expression of IGFBP-2 (insulin-like growth factor binding protein-2). The molecules of the invention are used to delay progression of hormone-regulated tumours, particularly of breast or prostate, to the hormone-independent state, to delay metastatic progression to the bone of IGF-1-sensitive cancers and to treat hormone-responsive cancers by inducing apoptosis, after hormonal withdrawal. The present nucleic acid sequence represents one of a collection (ABR70313-ABR70375) of antisense IGFBP-2-oligodeoxynucleotides (ODN) that were used in the invention for prostate and other endocrine tumour therapy
                                                                                                                                                                                                                                                                                                                                                                                                T cell receptor; TCR; MHC; major histocompatibility complex; cancer; imaging; lung tumour; autoimmune disease; neoplastic disease; cytostatic; immunosuppressive; virucide; antibacterial; protozoacide; yeast; mutant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New mutant soluble T cell receptor, useful e.g. for treating neoplasia or autoimmune disease, has higher affinity for ligand than the wild type.
                                                                                                                                                                                                       Gaps
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                                                                                                                                                                             Query Match 0.4%; Score 15.2; DB 1; Length 21; Best Local Similarity 85.0%; Pred. No. 1.6e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /product= "VbetaCDR3 mutant peptide"
/note= "No start and stop codon"
/partial
                                                                                                                                                          Sequence 21 BP; 1 A; 8 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                             Yeast VbetaCDR3 mutant peptide DNA #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                            2504 CCACCCGGTGGCACCCGTG 2523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7; Page 13; 36pp; English.
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Claim 3; Page 13; 36pp; English.
                                                                                                                                                                                                                                         CCATCCGGGGGGACCCCGAG 1
                                                                                                                                                                                                                                                                                                         AAD39797 standard; DNA; 21 BP
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/*tag= a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Saccharomyces cerevisiae.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KRAN/) KRANZ D M.
(WITT/) WITTRUP K D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-471080/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HOLL/) HOLLER P D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB; AAE24738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2002058253-A1.
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affinity for a ligand than wild-type T cell receptors (TCRs). TCRs are used, generally in the same way as antibodies, to identify ligands, to bind a selected peptide/MHC (major histocompatibility complex) ligand, as diagnostic probes for specific peptide/MHC molecules on cell surfaces, e.g. for diagnosing particular types of cancer or, in imaging, to locate and estimate size/number of tumours, also for diagnosis of many neoplastic diseases, to block autoimmune destruction of cells, to treat protozoa) T cells that include a gene for their receptors are used for recognition of selected peptide/MHC bearing cells. The present sequence is a yeast VbetaCDR3 (complementarity determining region) mutant peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an insulin regulator construct comprising a glucose response element (GIRE) of a liver pyruvate kinase (LPK) gene promoter and an insulin-sensitive element (ISE) of an insulin-like growth factor binding protein-1 (IGFBP-1) basal promoter. The construct is used to treat or prevent diabetic complications, to regulate insulin production, to modulate hyperglycaemia (without severe hypelypagmenia), to increase fat catabolism and creazed protein catabolism. The construct provides stimulation of insulin expression by glucose (but not other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Glucose response element; GIRE; liver pyruvate kinase; LPK; ss; IGFBP-1; insulin-sensitive element; ISE; basal promoter; hyperglycaemia; insulin; insulin-like growth factor binding protein-1; hypoglycaemia; glucose; fat catabolism; protein catabolism; carbohydrate; glucocorticoid; human; glucagon; euglycaemia; diabetes; fasting; ketogenesis; ketoacidosis; phepatocyte; heptoma; cellular protein degradation; antidiabetic; primer; intracellular hormone receptor; insulin regulator construct; anabolic; reverse transcriptase; RT-PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New insulin regulator cassette, useful e.g. for treating diabetes, provides specific, glucose-inducible transgenic expression of insulin in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 21 BP; 2 A; 4 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3203 TGGAGGATCCCTTCCAAGCC 3222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           В.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-OCT-2001; 2001US-00972916.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.4%;
Best Local Similarity 85.0%;
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2002107198-A1.
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Example 10; Page 119; 714pp; English.

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carbohydrates) and glucocorticoids and inhibition by glucagon so that a combination of these effects can maintain nearly euglycaemic conditions in diabetics during short-term feating, large carbohydrate loads or when fed ad libitum and prevent pathological ketogenesis and ketoacidosis, thus inhibiting the long-term complications of diabetes. The properties of the construct are essentially specific for hepatocytes and well-differentiated hepatoma lines and insulin expression in these cells may have effects additional to those provided by secreted insulin, e.g. inhibition of cellular protein degradation, and inhibition, or stimulation of other intracellular hormone receptors. This sequence represents a reverse transcriptuse PCR (RT-PCR) primer used to amplify human insulin, used in the scope of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 3 A; 7 C; 7 G; 4 T; 0 U; 0 Other;
          886666666666888
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0.4%; Score 15.2; DB 1; Length 21; 35.0%; Pred. No. 1.6e+03; 3; Indels 0; Mismatches 1611 GTGCATCCACAGGGACCTGG 1630 GCGCATCCACAGGGCCATGG 2 82.0%; Local Similarity 85.0 nes 17; Conservative 21 Query Match Matches à g

0

Gaps

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ABS97565 standard; DNA; 21 ABS97565;

(first entry) 23-DEC-2002

Human epoxide hydroylase 2 polymorphic sequence #56.

W Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
Cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;
Cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;
Adrenegic receptor hubbar translocator; ARNT; cathepsin S; CTSS;
TY1 hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
Cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
W poxide hydroxylase 2; EPKZ; 5-11poxygenase activating protein; FLAPP;
Qultathione-S-transferase 12; GGT12; histamine-N-methyl transferase;
NMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase;
NMDP-glucuronosyl transferase 2; NQO2; sulfoctransferase thermolabile; STM;
UDP-glucuronosyl transferase 2; UGT2B15; urokinase receptor;
W UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor;
W multidrug resistance associated protein 3; cancer; prostate;
M multidrug resistance associated protein 3; cancer; prostate;
A cetylcholine muscarinic receptor; CHWR1; CHWR2; CHWR5;
A altered drug metabolism; cardiovascular function; colorectal tumour;
W single nucleotide polymorphism.

Homo sapiens

WO200257410-A2

25-JUL-2002

28-NOV-2001; 2001WO-US044838.

28-NOV-2000; 2000US-00724389.

(DNAS-) DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes eg. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traitis.

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this invention relates to the Sequence or an isolated nuclear action molecule comprising at least one base variation from that of a known controlled by the control of the comprising at least one base variation from that of a known correction receptor betail (ADBRI), cytochrome P450 02E1 (CYP45002E1), adrenatic receptor betail (ADBRI), aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (AANT), cathepsin S (CTSS), cyclooxegenses 2 (COX2), diazepam binding inhibitor (DBI), epoxide hydrocylase 2 (EPHX2), 5-lipoxygenase activating thibitor (DBI), epoxide hydrocylase 2 (EPHX2), 5-lipoxygenase activating transferase (HNMT), MADPH quinome oxidoreductase 2 (MQ2)) histamine-N-methyl transferase (HNMT), MADPH quinome oxidoreductase 2 (MQ2)) histamine-N-methyl transferase (UGT2BI), urokinase receptor (UPA), multidrug resistance 1 (UGT2BI), upp-glucuronosyl transferase 2 (MQ2), cyphan nuclear receptor (NRI2), or acetylcholine muscarinic (MRB1), orphan nuclear receptor (NRI2), or acetylcholine muscarinic creeptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR4 or CHMR4 or CHMR5) sequence. Creeptor 1, 2, 3, 4, or 5 (CHMR1), or acetylcholine muscarinic creeptor (Linkage markers for locating and characterising the genes that creeptor 1, 2, 3, 4, or 5 (CHMR1), creating the genes responsible for avariety of disorder-related creeptor (CT traits as a result of their e.g., overexpression, occitained in CTP4501A1, CTP4501A2, CTP4501A2, and/or treating the disorders. The nucleic acid molecules comprising the and/or MDR3 are useful for screening individuals for altered drug metabolism. The polymorphic sequences ontained in CTP4501A1, CTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                      This invention relates to the sequence of an isolated nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 4 A; 2 C; 13 G; 2 T; 0 U; 0 Other;
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Gaps ö 0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03; tive 0; Mismatches 3; Indels Query Match 0.4 Best Local Similarity 85.0 Matches 17, Conservative

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AAL43996 standard; DNA; 21 BP. AAL43996; RESULT 1437 AAL43996/

(first entry) 27-SEP-2002

Reproductive recombination virus-related oligonucleotide SEQ IN #16.

Gene therapy, ss, reproductive recombination virus, tumour cell killing; Epstein-Barr virus; cancer-suppressing gene; vascular inhibition gene; cell factor gene; prodrug-converting enzyme gene; cell death gene; nasopharyngeal cancer; Hodgkin's lymphoma; gastric cancer; PCR; primer.

WO200256917-A1.

25-JUL-2002.

17-JAN-2002; 2002WO-CN000025.

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                                                                                                                                                                                                           The invention comprises a reproductive recombination virus capable of killing specifically tumour cells associated with Epstein-Barr (EB) virus. The virus of the invention comprises an insertion of a target gene into the non-reproduction-requiring domain in the virus gene group (the target gene is a cancer-suppressing gene, vascular inhibition gene, cell factor gene, prodrug-converting enzyme gene or cell death gene). The reproductive recombination virus of the invention is useful for treating nasopharyngeal cancer, Hodgkin's lymmhomma and gastric cancer. The present DNA sequence was used in an example of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New PR Family Member 2 oligonucleotide, useful for preparing a composition for modulating cell growth for treating cancer or diseases of cell degeneration, e.g., Alzheimer's disease or insulin-dependent diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, PR Family Member 2; PFM2; PFM PR domain, PFM zinc finger domain, PFM ZF domain, modulation of cell growth; cancer; cell degeneration disease, Alzheimer's disease, Parkinson's disease, insulin-dependent diabetes mellitus; IDDM; neuroprotective; antidiabetic; cytostatic; RT-PCR;
                                                                                                                         Construction of reproductive recombination virus able to specifically kill Epstein-Barr associated tumor cells, useful in drugs for treating e.g. nasopharyngeal cancer, Hodgkin's lymphoma and gastric cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                   Score 15.2; DB 1; Length 21;
Pred. No. 1.6e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 9 A; 6 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reverse transcriptase-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                             1457 GTAACCTGCGGGAGTTTCTG 1476
                                        (VIRG-) VIRGENE BIOTECHNOLOGY LTD
                                                                                                                                                                                  Example 2; Page 13; 44pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PFM2 cDNA RT-PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 Gracciecrecarrirere 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA08083 standard; DNA; 21 BP
              18-JAN-2001; 2001CN-00105247.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00389956.
                                                                                                                                                                                                                                                                                                                                                                                                   85.0%;
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 85.0'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-2003 (first entry)
                                                                    Che X, Sham S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BURN-) BURNHAM INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-669568/63.
                                                                                                WPI; 2002-566772/60.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Qian Q,
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                                        The present invention relates to the isolation of human and mouse PR Family Member (PFM) proteins, and the polymucleotide sequences encoding them. Also disclosed are PFM PR and PFM zinc finger (ZF) domains, and the polymucleotide sequences encoding them. The invention also discloses PFM oligonucleotides and methods for detecting a PFM polymucleotide sequence in a sample. The PFM polypeptide and polymucleotide sequence for preparing a composition for modulating cell growth for treating cancer or diseases of cell degeneration, e.g. as Alzheimer's disease, Parkinson's disease or insulin-dependent diabetes mellitus (IDPM). The
                                                                                                                                                                                                                                                             present sequence represents a primer used in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting the presence of a target nucleic acid sequence on a sample nucleic acid strand, useful for human leukocyte antigen tissue typing, comprises contacting a sample with a diagnostic probe under hybridizing
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                         Seguence 21 BP; 1 A; 6 C; 5 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Probe sequence #36 related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human leukocyte antigen; HLA; probe; PCR; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1722 GAAGACAACCAACGCCGGC 1741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 32; 62pp; English.
Example 2; Col 29; 95pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 GAAGACAATCAACAGCGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB78533 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-363216/34.
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                                                                                                                                                                                                                                                                                           invention
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Human; ss; PCTR5; colorectal cancer; adenomatous polyposis coli;

Wyelogenous leukaemia; congenital neonatal autoimmune thrombocytopenia;

Wovarian cancer; brain tumour; breast cancer; glioma; astrocytoma;

W renal cell carcinoma; melanoma; clear cell carcinoma;

Granular cell carcinoma; neurological disorder;

W neurodegenerative disorder; nerve trauma;

M familial myelodysplastic syndrome; Charcot-Marie-Tooth neuropathy;

Gardner syndrome; mental health condition; immunological disorder;

M allergy; asthma; lung disease; reproductive disorder; deafness;

M glycoprotein deficiency; desmoid tumour; turcot syndrome;

N plastos; schistosoma mansoni infection; spinocerebellar ataxia;

M plasmodium falciparum infection; groenow's corneal dystrophy;

M lattice corneal dystrophy; probe.
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                                                                                            Gaps
and does so at a lower cost. The present sequence represents a probe
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                                                               0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
tive 0; Mismatches 3; Indels
                                       Sequence 21 BP; 5 A; 5 C; 10 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Herrmann J,
Rastelli L;
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Mezes PS,
                                                                                                                  231 CTGGACACGCCCCGAGCGGA 250
                                                                                                                                            credadadaccccaaccea 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAR-2001; 2001US-00800198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-MAR-2000; 2000US-0186592P
                                                                                                                                                                                                           ADB32049 standard; DNA; 21
                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ф Е,
                                                                                       17; Conservative
                                                                                                                                                                                                                                                                                      Human FCTR5 cDNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (VERM/) VERMET C.
(FERN/) FERNANDES E.
(SHIM) SHINKETS R.
(HERK/) HERRMANN J.
(MAJU/) MAJUMDER K.
(MAJU/) MAJUMDER K.
(MISH/) MISHRA V.
(MISH/) MISHRA V.
(MEZK/) MISHS P.S.
(MASK/) MASTELLI I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fernandes E
J, Mishra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-625633/59
                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003087816-A1.
                the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Macdougall J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                              04-DEC-2003
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                                                                Query Match
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                                                                              Local
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                                                                                        Matches
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New FCTRX polypeptide and encoding polynucleotide, useful for preventing or treating FCTRX-related disorders, such as cancer, autoimmune, neurodegenerative, gastrointestinal, reproductive and inflammatory

The invention relates to FCTRX polypeptides and the polynucleotides encoding them. The sequences of the invention are useful for the

Example 2; Page 146; 155pp; English.

diseases.

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manufacture of a medicament for diagnosing and treating disorders associated with the FCTRX polypeptide, such as colorectal cancer, adenomatous polyposis coli, wpelogenous leukaemia, congenital neonatal autoimmune thrombocytopenia, malignant ovarian tumours, malignant brain tumours, mammary tumours, malignant ovarian tumours, malignant brain carcinoma, breast adenocarcinoma, ovarian carcinoma, melanomas, clear cell and granular cell carcinoma, novarian carcinoma, melanomas, clear cell and granular cell carcinoma, nevelogical disorders, contrological disorders, nevel trauma, familial myelodysplastic syndrome, Charcot Marie-Tooth neuropathy, Gardner syndrome, mental health conditions, immunological disorders, allergy and infection, asthma, lung diseases, male and female reproductive disorders, dasfness, glycoprotein deficiency, desmoid tumour, turcot syndrome, liver cirrhosis, hepatitis correction, spinocerebellar ataxia, plasmodium falciparum infection, infection, spinocerebellar ataxia, plasmodium falciparum infection, crepresents a probe used to detect cDNA encoding an FCTRX polypeptide of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises a human antibody for inhibiting platelet aggregation by its exclusive binding to the activated state of platelet integrin receptor GPIDA/IIIa. The antibody of the invention is useful for preparing a diagnostic composition for determining the number of activated thrombocytes or for blocking the platelet integrin receptor on thrombocytes. The antibody of the invention is useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antibody, useful for preparing a composition for determining the number of activated thrombocytes or for blocking the platelet integrin receptor on thrombocytes for treating e.g., thrombosis or myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody; platelet aggregation inhibition; platelet integrin receptor; GPIIb/IIIa; activated thrombocyte; thrombosis; myocardial infarction; primer; 88; human; PCR.
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                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 3 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schwarz M, Knackmuss S, Peter K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunoglobulin heavy chain PCR primer #9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1182 GGCCCGGCTGACCCTGGGCA 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 GGCCCAGCTGACCCTGCTCA 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE03295 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 85.0
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              infarction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE03295;
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double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epliepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; TGFB1.

TGFB1-targeted sinA DNA-RNA hybrid - SEQ ID 882.

(first entry)

26-FEB-2004

ADG30316;

ADG30316 standard; RNA; 21 BP.

ADG30316,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polynucleotide useful for detecting single nucleotide polymorphisms
thrombosis or myocardial infarction. The present DNA sequence represents a PCR primer that was used in an example of the invention.
                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human; single nucleotide polymorphism; microarray; side effect; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism detection primer, SEQ ID No 1849.
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                                                                                                                                                                                 ch 0.4%; Score 15.2; DB 1; Length 21; 1 Similarity 85.0%; Pred. No. 1.6e+03; 17; Conservative 0; Mismatches 3; Indels
                                                                                                              Sequence 21 BP; 3 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 9 A; 9 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; SEQ ID NO 1849; 704pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detection method of the invention.
                                                                                                                                                                                                                                                                                                                                           854 AGGAGGAGCTGGTGGAGGCT 873
                                                                                                                                                                                                                                                                                                                                                                                  ADF88266 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-FEB-2002; 2002JP-00034717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-FEB-2002; 2002JP-00034717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-820454/77.
                                                                                                                                                                                                                             Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP2003235571-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in human gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF88266;
                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1442
ADP88266/C
XX
AC ADP88266/C
XX
AC ADP88266
DT 26-FBB-2
XX
AC ADP88266
DT 26-FBB-2
XX
AC ADP88266
DX Synthem:
OS Synthem:
OS Synthem:
OS Synthem:
OS Homo saj
XX
AC ADP88266
DX AC AUG-:
CC AUG-:
DX AC AUG-:
DX AC AUG-:
CC AUG-:
DX AC AU
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Fosnaugh K;

Pavco P,

Chowrira B,

Beigelman L,

(SIRN-) SIRNA THERAPEUTICS INC

11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-03867B2P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SER-2002; 2002US-040929P.

20-FEB-2003; 2003WO-US005028

WO2003074654-A2.

12-SEP-2003

Unidentified.

Synthetic.

Thompson J;

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(sinA) molecule that down-regulates expression of an endogenous mammaliantarget gene comprising one or more chemical modifications and each strand of the double-stranded sinA comprises about 21 nucleotides. The sinA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, dissue or organism. The disease or condition may include pulmonary disease such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a double-stranded short interfering nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 4 A; 3 C; 8 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 882; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1595 ACTTGGCCTCCCAGAAGTGC 1614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 ACTCTGCCTCCCAAAGTGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 85.09
Matches 17, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J, Belyeam... R. Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activity
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Gaps

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3; Indels

0; Mismatches

17; Conservative Local Similarity

Query Match

Best Loca Matches

TGTGCGTGTGTGTGTGTG 2349

2330 20

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rerecerciererererere 1

0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03;

Antisense oligonucleotide; RNase; cancer; B cell lymphoma-associated gene; BCL2; STAT3; HER-2; focal adhesion kinase; FAK; melanoma; inflammatory disease; tumour necrosis factor alpha; TNF-alpha; probe; ss; human.

08-MAY-2002; 2002US-00142666. 18-JUL-2001; 2001US-0306229P.

US2003171315-A1. Homo sapiens.

11-SEP-2003.

Brown BD, Riley TA; WPI; 2003-898297/82.

(BROW/) BROWN B D. (RILE/) RILEY T A.

Human antisense oligonucleotide #5.

(first entry)

03-JUN-2004

ADM41184;

ADM41184 standard; DNA; 21 BP.

1445

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the instraion codon, coding regalon, 5' or 3' end genomic flanking regions, c' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entill ammatory steroid in a subject, for reducing levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed are for interval and the printed are formal and the printed and the formal and the formal directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                               antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aguilar D;
                                                                                                                                                                                                                Human; antisense; lung dysfunction; nasal airway dysfunction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1; Length 21; Pred. No. 1.6e+03; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 0 A; 6 C; 14 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 11555; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Katz E,
                                                                                                                                                                   Human C/EBP antisense fragment no.2173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sandrasagra A, Ka
, Shahabuddin S;
                                            ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%;
                                       ABZ96313 standard; DNA; 21
                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 85.09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Li Y, Sar
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                     WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                           17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                               31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Miller S,
                                                                                 ABZ96313;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nyce JW,
RESULT 1444
                   ABZ96313
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or

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The invention relates to an antisense oligonucleotide comprising a domain that recruits an RNase and inhibits the function of a gene associated with a human disease, where the improvement comprises incorporation of two juxtaposed universal bases in the oligonucleotide. The antisense oligonucleotides are used for treating cancer associated with an overexpression of the B cell lymphoma-associated gene (BCL2), STAT3, HER-and focal adhesion kinase (PAK), melanoma and inflammatory diseases associated with an expression of tumour necrosis factor (TNF-alpha), and for inhibiting function of a gene associated with a human disease. This sequence represents a human antisense oligonucleotide used in the scope
                                                                                                                                                                                                                                                                                                                                                                                                                                     comprises at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                  Improved antisense oligonucleotide for treating e.g. cancer, least two juxtaposed universal bases in the oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 15.2; DB 1; Length 21; Best Local Similarity 70.0%; Pred. No. 1.6e+03; Matches 14; Conservative 3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 3 A; 6 C; 6 G; 1 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; SEQ ID NO 13; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1434 GCTGGTGGAGTACGCGGCCA 1453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human C/EBPN DNA fragment 2173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GCUGGUUGAGUACUCCGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABD20222 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABD20222;
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Gaps

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1476 GCGGGCGCGGCCCCCGG 1495

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Penicillium citrinum reductase mutagenesis PCR primer SEQ ID NO:13.

(first entry)

22-APR-2004

ADH44226;

ADH44226 standard; DNA; 21 BP.

RESULT 1447

reductase; Penicillium citrinum; enzyme; (S)-4-halo-3-oxobutyrate ester; modified reductase; beta-keto acid reduction; organic synthesis;

mutagenesis; PCR primer; ss

Penicillium citrinum.

Synthetic

3P1386961-A2.

04-FEB-2004.

30-JUN-2003; 2003EP-00254142. 03-JUL-2002; 2002JP-00194344. SUMO) SUMITOMO CHEM CO LID

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiniflammatory, antiathmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or creating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The composition or a vailability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The cultinammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated conflammation, allergies, asthma, independent engine content of the anti-sense oligos corresponding to the main frame into the target RNA serves to prevent the breakdown of the charget parameter of the midles present in the target RNA serves to prevent the breakdown of the conflammation and or prevent the breakdown of the particular and or prevent the present in the target RNA serves to prevent in the carget RNA serves to prevent in the carget RNA serves to prev
                                                                              beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ds.
respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D;
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 21 BP; 0 A; 6 C; 14 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; SEQ ID NO 11555; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           into products that
eart, kidney, etc, t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                               23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the oligonucleotides into e.g., lung, brain, heart,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                      WO200285309-A2
                                                                                                                                                                                                                                                                                                                           31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce JW, 1
Miller S,
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New modified reductases comprising an amino acid sequence having a substitution at position 54 and/or 104, or a further amino acid deletion, substitution or addition, useful for reduction reaction of e.g. beta-keto

WPI; 2004-135408/14.

Shimizu M;

Asako H,

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The present invention describes a modified reductase (I) isolated from Penicillium citrinum comprising: (a) a sequence of 325 amino acids (SEQ CC ID NO: 1, ADH44214) having a substitution at amino acid position 54 and/or 104; or (b) an amino acid sequence of (a) having further deletion, and/or 104; or (b) an amino acid sequence encoding the amino acid sequence encoding the amino acid sequence encoding the amino acid sequence of the reductase defined above; (2) a vector comprising the polynucleotide; (3) a transformant comprising the polynucleotide or vector; (4) producing (S)-4-halo-3-hydroxybutyrate ester by reacting 4-compression (S) material; (5) modified (S)-4-halo-3-hydroxybutyrate ester by reacting 4-complexity (B) modified enzyme gene by substituting at least one single material; (5) modified enzyme gene by replacing at codon that corresponds to producing a modified enzyme gene by replacing a codon that corresponds to at least one of the amino acide of the positions 54 and 104 of SEQ ID NO: 1 with a codon that corresponds to another amino acid(s) in a nucleotide sequence that encodes SEQ ID NO: 1. The modified reductase can be used for reduction reaction, specifically reduction reaction of beta-keto acid, and in organic synthesis reaction for the production of compounds used as active ingredients of medicaments, agrochemicals or their intermediates, especially optically active compounds. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; SEQ ID NO 13; 40pp; English.
tissue environment and thereby, to
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Gaps
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                                                            Length 21;
                                                                                           Indels
                              Sequence 21 BP; 6 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                            0.4%; Score 15.2; DB 1;
85.0%; Pred. No. 1.6e+03;
                                                                                              0; Mismatches
in an example from the present invention.
                                                                                                                               2443 TGGTGCTGCAGCACCGAGGG 2462
                                                                                                                                                           1 recraciaciaciaciacia 20
                                                                            Local Similarity 85.0 es 17; Conservative
                                                              Query Match
                                                                                             Matches
                                                                          's Best
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Gaps

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Indels

0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03;

0; Mismatches

17; Conservative

Local Similarity

Query Match Best Loca Matches 1476 GCGGGCGCGCGCCCCGG 1495 2 ściegesciesciesciercz

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The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; probe; NOVX; autoimmune disease; Alzheimer's disease; stroke; allergy; Parkinson's disease; multiple sclerosis; anxiety; pain; diabetes; graft Hersus host disease; pancreatitis; obesity; uloer; anaemia; cancer; viral infection; bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                              Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.4%; Score 15.2; DB 1; Length 21; Best Local Similarity 85.0%; Pred. No. 1.6e+03; Matches 17; Conservative 0; Mismatches 3; Indels
                                           human; single nucleotide polymorphism; SNP; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 8 A; 9 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; SEQ ID NO 3561; 2627pp; Japanese
                                                                                                                                                                                                                                                                                                       (KAGA-) KAGAKU GLJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotide polymorphism in human gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human NOVX-associated primer/probe #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2323 GTGTGTGTGTGTGTGT 2342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , 2001US-0310291P.
; 2001US-0310291P.
; 2001US-0311292P.
; 2001US-0311379P.
; 2001US-0311379P.
; 2001US-0313201P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 GAGTGTGTGCACGTGTGTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADJ51062 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                    08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-AUG-2002; 2002US-00210281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      parasitic infection; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17; Conservative
                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-093977/10.
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                                                                                                                                JP2003259875-A.
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14-AUG-2001;
17-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-AUG-2001;
03-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-AUG-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-AUG-2001;
                                                                                                                                                                         16-SEP-2003
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                                                                                      Synthetic.
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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New PR family member nucleic acid and polypeptide, useful for diagnosing, preventing and treating proliferative disorders, such as cancer, scleroderma, arthritis or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention provides novel PR family member (PFM) polypeptides and polynucleotides. These sequences are characterised by encoding or containing a PR domain and a zinc finger (ZF) domain. The PFM nucleic acid molecules, oligonucleotides, polypeptides and peptides are useful for modulating cell proliferation, diagnosing, preventing and treating proliferative disorders (e.g. cancer, scleroderma, arthritis, alcoholic liver cirrhosis, keloid and hypertrophic scarring or atherosclerosis) or disorders of cell degeneration (e.g. Duchenne's muscular dystrophy, insulin-dependent diabetes mellitus, Parkinson's disease, Huntington's disease, Alzheimer's disease or paralysis). PFM nucleic acid molecules are useful as probes and primers to detect PFM nucleic acid molecules are useful as probes useful in gene therapy. The present sequence is human PFM DNA amplifying PCR primer.
                                                                                                                                                                                                                                                           arthritis, alcoholic liver cirrhosis, keloid, hypertrophic scarring, atherosclerosis, cell degeneration disorder; Duchenne's muscular dystrophy; insulin-dependent diabetes mellitus, Parkkinson's disease, Huntington's disease, Alzheimer's disease; paralysis, gene therapy; human; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                         family member; PFM; proliferative disorder; cancer; scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 21 BP; 1 A; 6 C; 5 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                             Human PFM2 DNA amplifying PCR primer, PFM2.18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; SEQ ID NO 82; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1722 GAAGACAACCAACGCCGGC 1741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 GAAGACAATCAACAGCGGGC
                                                             ADJ32263 standard; DNA; 21 BP
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                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BURN-) BURNHAM INST
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                                                                                                                                                                                                                                                                                                                                                                                                                                         US2003236395-A1.
                                                                                                                                                  22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-SEP-1999;
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                                                                                                         ADJ32263;
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                    RESULT 1448
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Matches

à 요 RESULT 14.
ADK94532//
ID ADK9.
XX
AC ADK9.
XX
DT 06-M

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Gaps

Seguence 21 BP; 3 A; 4 C; 6 G; 8 T; 0 U; 0 Other;

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New NOVX polypeptides and polynucleotides, useful in diagnosing, treating
                                                                                                                                                                                                                                               or preventing diseases or conditions, e.g. autoimmune disease,
Alzheimer's disease, diabetes, graft versus host disease, cancer or viral
or bacterial infections.
                                                                                                                                                                                                                                                                                        The invention relates to an isolated NOVX polypeptide (of 44 disclosed)
                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 127; 342pp; English.
      21-AUG-2001; 2001US-0314031P.
23-AUG-2001; 2001US-0314466P.
28-AUG-2001; 2001US-0315403P.
29-AUG-2001; 2001US-0315853P.
05-MAR-2002; 2002US-036175P.
05-MAR-2002; 2002US-0361735P.
                                                     GORMAN L.
ZERHUSEN B D.
EDINGER S R.
PADIGARU M.
                                                                                                                                                                                                                            WPI; 2004-168942/16.
                                                                                                                                                                                                        Stone
                                                                                                                                                                                          Gorman L,
                                                                                                                                                                                                         Sciore P
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                                                                                            (ZHON/)
(PATT/)
(MILL/)
                                                                                                                       (PENA/)
(BURG/)
(SCIO/)
(STON/)
(TAUP/)
                                                                                                                                                               (ROTH/)
                                                     GORM/)
                                                                                GUOX/)
                                                                                                                                                         CASM/)
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                                                                                       KEKU/)
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ADJ97593
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EA, Burgess CE;
                                                                                                                                                                                                                                                     Ji W, Pena CEA, Burges
Casman SJ, Rothenberg ME;
                                                                                                                                                                                                                                     Zerhusen BD, Edinger SR, Padigaru M,
Patturajan M, Miller CE, Ji W, Pena C
                                                                                                                                                                                                                                                                    Taupier RJ,
                                                                                                                                                                                                                                                                                 Malyankar UM, Boldog FL;
                                                                                                                                                                       ROTHENBERG M E. MALYANKAR U M.
                                                                                                                                                                                                                                                     Patturajan M,
              ZHONG M.
PATTURAJAN M.
MILLER C E.
                                                                                                                          STONE D J.
TAUPIER R J.
CASMAN S J.
                                                                          PENA C E A.
BURGESS C E.
SCIORE P.
                                                                                                                                                                                                         BOLDOG F L.
KEKUDA R.
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Disclosure; SEQ ID NO 366; 218pp; English. retinopathy and cancer. WPI; 2004-203472/19. The invention fraintes to an isolated who purpoperate to the invention fraintes to an isolated having at least 95% sequence comprising the mature form, a sequence having at least 95% sequence identity to NOVX or a sequence comprising one or more conservative as ubstitutions in the amino acid sequence of NOVX. Also included are a composition comprising who acrier, a kit comprising, in one or more containers, the composition, a method of identifying an agent that binds to NOVX, a method for identifying a potential therapeutic agent for use in treatment of a pathology related to aberrant expression or aberrant physiological interactions of NOVX, a method for screening for a modulator of activity of or of latency or predisposition to a pathology associated with NOVX, a method for modulating the activity of NOVX, a method for modulating the nucleic acid molecule a cell comprising the vector comprising the nucleic acid molecule, a cell comprising the vector, an antibody that immunospecifically binds to NOVX, a method of producing NOVX (comprising culturing the presence of or predisposition to a disease associated with altered levels of expression of NOVX or the nucleic acid molecule in a sample, a method for determining the presence or amount of NOVX or the under conditions that lead to expression of the polypeptides or expression of NOVX or the polypeptides cueful in the manufacture of a medicament for treating a syndrome conditions that lead to expression of the polypeptides cueful in the manufacture of a medicament for treating a syndrome associated with a human disease associated with NOVX. The polypeptides and nucleic acid molecules are useful in diagnosing, treating or all presence or and nucleic acid molecules are useful in diagnosing, treating or all presence or and nucleic acid molecules are useful in diagnosing the polypeptides. preventing diseases or conditions, e.g. autoimmune disease, Alzheimer's disease, stroke, allergies, Parkinson's disease, Huntington's diseases, multiple sclerosis, anxiety, pain, diabetes, graft versus host disease, pancreatitis, obesity, ulcers, anaemia, cancer, viral or bacterial and parasitic infections (many more diseases and disorders are listed in the specification). The present sequence is a primer or probe included in the sequence listing but not mentioned anywhere else in the specification.

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This invention relates to novel compositions that comprise short interfering RNA (siRNA) molecules, which can be used to inhibit can dendicensis. Specifically, it refers to siRNAs that target and cause RNAi-induced degradation of mRNA from human vascular endothelial growth factor (WEGF), the VEGF receptor (Flt-1) and the Flk-1/KDR (kinase domain cegion) genes; as well as mutants derived thereof. The present invention describes sense and antisense RNA strands that form an RNA duplex and cherraped of the target mRNA, such that expression is inhibited and the target degraded. As such, siRNA administered in combination with a therapeutic gent is useful for treating diseases associated with analogenesis and the overexpression of VEGF, which include diabetic retinopathy, agenciated macular degeneration, inflammatory disease, psoriasis and reherapeutic related macular degeneration, inflammatory disease, psoriasis and reherapeutic cancers including breast, retinoblatema, wilm's tumour and lymphoma. Accordingly, these compositions exhibit cytostatic, antidiabetic, ophthalmological, antiinflammatory, antipsoriatic, antidiabetic, antidiabetic antiarthritic activities. This oligonucleotide is a human Flt-1 DNA coligo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               age-related macular degeneration; inflammatory disease; psoriasis; rheumatoid arthritis; cancer; breast; retinoblastoma; Wilm's tumour; lymphoma; cytostatic; antidiabetic; ophthalmological; antiinflammatory; antipsoriatic; antirheumatic; antiarthritic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel short interfering RNA (siRNA) comprises sense and antisense RNA strands, useful for inhibiting expression of human vascular endothelial growth factor mRNA, for treating angiogenic disease, e.g. diabetic
                                              Gaps
                                                                                                                                                                                                                                                                                                                                                              Human Flt-1 DNA sequence, a target for siRNA inhibition SeqID 366.
                                                                                                                                                                                                                                                                                                                                                                                                  human, ss; short interfering RNA, siRNA, angiogenesis;
vascular endothelial growth factor; VEGF, VEGF receptor; Flt-1;
Flk-1/KDR; kinase domain region; diabetic retinopathy;
                                           ;
0
0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
iive 0; Mismatches 3; Indels
                                                                                   3669 CATGGCTCAGGGTGGTCTCT 3688
                                                                                                                             caregercagarerrerer 21
                                                                                                                                                                                                                                    ADJ97593 standard; DNA; 21 BP
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14-NOV-2002; 2002US-00294228.
                                                                                                                                                                                                                                                                                                                       06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UYPE-) UNIV PENNSYLVANIA.
                        Best Local Similarity 85.0
Matches 17, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tolentino MJ, Reich SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004009769-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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      Query Match
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vascular endothelial growth factor; VEGF; VEGF receptor; Flt-1; vascular endothelial growth factor; VEGF; VEGF receptor; Flt-1; Flk-1/KDR; kinase domain region; diabetic retinopathy; age-related macular degeneration; inflammatory disease; psoriasis; rheumatoid arthritis; cancer; breast; retinoblastoma; Wilm's tumour; lymphoma; cytosteatic; antidiabetic; ophthalmological; antiinflammatory; antipsoriatic; antirheumatic; antiarthritic.
                                             Gaps
                                                                                                                                                                                                           Human Flt-1 DNA sequence, a target for siRNA inhibition SeqID 414
                                            ;
0
                      Score 15.2; DB 1; Length 21;
Pred. No. 1.6e+03;
                                            3; Indels
                                                                                                                                                                                                                                human; ss; short interfering RNA; siRNA; angiogenesis;
Sequence 21 BP; 7 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                            0; Mismatches
                                                               1300 ATGCTGAAAGACGATGCCAC 1319
                                                                                     Argercaaagagggggcgc 21
                                                                                                                                           BP
                      0.4%;
                                                                                                                                                                                                                                                                                                                                                                                  18-JUL-2003; 2003WO-US022444.
                                                                                                                                                                                                                                                                                                                                                                                                        24-JUL-2002; 2002US-0398417P
14-NOV-2002; 2002US-00294228
                                                                                                                                          ADJ97641 standard; DNA; 21
                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYPE-) UNIV PENNSYLVANIA.
                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tolentino MJ, Reich SJ;
                     Query Match
Best Local Similarity
Matches 17; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-203472/19.
                                                                                                                                                                                                                                                                                                                                         WO2004009769-A2.
                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                     06-MAY-2004
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                                                                                                                                                               ADJ97641;
                                                                                                                       RESULT 1452
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interfering RNA (sirRNA) molecules, which can be used to inhibit angiogenesis. Specifically, it refers to siRNAs that target and cause angiogenesis. Specifically, it refers to siRNAs that target and cause RNA-induced degradation of mRNA from human vascular endothelial growth factor (VEGP), the VEGP receptor (FL-1) and the FIR-1/KDR (Kinase domain region) genes, as well as mutants derived thereof. The present invention describes sense and antisense RNA strands that form an RNA duplex and bind to the target mRNA, such that expression is inhibited and the target degraded. As such, siRNA administered in combination with a therapeutic ageraded. As such, siRNA administered in combination with a therapeutic degraded. As such, retaining diseases associated with angiogenesis and the overexpression of VEGF, which include diabetic retinopathy, age-related macular degeneration, inflammatory disease, psoriasis and cancers including breast, retinoblastoma, Wilm's tumour and lymphoma. Accordingly, these compositions exhibit cytostatic, antidiabetic, ophthalmological, antiinflammatory, antipsoriatic, antirheumatic and antiarthritic activities. This oligonucleotide is a human Flt-1 DNA coligo, a target for siRNA inhibition of the invention. This invention relates to novel compositions that comprise short

Novel short interfering RNA (siRNA) comprises sense and antisense RNA strands, useful for inhibiting expression of human vascular endothelial growth factor mRNA, for treating angiogenic disease, e.g. diabetic

Disclosure; SEQ ID NO 414; 218pp; English.

retinopathy and cancer.

Sequence 21 BP; 6 A; 5 C; 5 G; 5 T; 0 U; 0 Other;

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The present invention relates to a method for identifying a BBB (bloodbrain barrier)-specific protein, or its fragment, in endothelial cells of brain barriers (brain microvessel endothelial cells, BMEC). The method comprises conventional pre-purification of BMEC, freshly isolated from the brain by enzymatic digestion, treating the digest with a lysis buffer that destroys erythrocytes and apoptotic cells but retains at least 70% or EMEC in viable form, optionally further purification of the product, preparing subtractive cDNA libraries from BMEC and a subtractive tissue, cerifying clones from the subtracted cDNA banks with respect to expression, completing BBB-specific cDNA clones, and comparing expression cerifying clones from the subtracted cDNA clones, which are useful for identify BBB proteins or their fragments. The method is useful for identifying BBB-specific proteins which are useful for transporting substances across the blood-brain barrier and for treatment and diagnosis of diseases (none indicated) associated with dysfunction of this barrier. The method is simple, is performed under mild conditions and provides or machinging in haring that are produced, predeminantly consorting the analysiucus is dentification of proteins that are produced, predeminantly consorting the procession of proteins that are produced, predeminantly consorting the procession in the procession of proteins that are produced, predeminantly consorting the procession of proteins that are produced, predeminantly consorting the procession of proteins that are produced, predeminantly consorting the procession of proteins that are produced, predeminantly consorting the procession of the procession of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying blood-brain barrier protein, useful e.g. as drug transporter and for treatment or diagnosis, by subtractive hybridization using cDNA from brain capillary cells.
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d are specific
d in the
                                                          Gaps
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0.4%; Score 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03; ive 0; Mismatches 3; Indels
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85.0%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blood-brain barrier-specific protein; BBB; endothelial cellbrain capillary; brain microvessel endothelial cell; BMEC; substance transport; TKA-1; human; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wolf S, Jaeger M, Bangsow T, Bangsow C, Jordan D,
Oppolzer T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ESPL-) ESPLORA GMBH C/O TU DARMSTADT INST BIOCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human TKA-1 cDNA, primer ssTKA-1ctg.as1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; SEQ ID NO 31; 63pp; German.
                                                                                                           1609 AAGTGCATCCACAGGGACCT 1628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for the BBB. The present sequence examples of the present invention.
                                                                                                                                                             2 AAGTGCATTCATCGGGACCT 21
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                                                                                                                                                                                                                                                                                               ADO60532 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
  Query Match 0.4
Best Local Similarity 85.0
Matches 17; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                             RESULT 1453
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This invention relates to a novel isolated polynucleotide comprising a nucleic acid sequence, the human MRCKI gene located at position 11q13, and the novel human protein Kinase MRCKI gene located by it. The sequence of the invention has sequence homology to rat myotonic dystrophy kinase. related Cdc42 binding kinase (MRCK). The invention may be useful for diagnosing, prognosing, and treating kinase-related diseases, preferably diseases associated with aberrant expression of MRCKI. The present sequence is that of a short inhibitory (siRNA) sequence which is targeted equence data he human MRCKI gene and which is related to the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from MIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New human protein kinase MRCK1 with 65% sequence homology to rat myotonic dystrophy kinase-related Cdc42-binding kinase (MRCK), useful in diagnostics and as a drug target.
                                                                                                                                  protein kinase; MRCK1; myotonic dystrophy kinase; Cdc42 binding kinase; MRCK; kinase-related disease; short inhibitory RNA; siRNA; human; ss.
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65.0%; Pred. No. 1.6e+03;
tive 4; Mismatches 3; Indels
                                                               Human MRCK1 sense strand siRNA sequence SeqID161.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 4 A; 5 C; 5 G; 0 T; 7 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 161; 92pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                          07-NOV-2003; 2003WO-US035609.
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09-SEP-2004 (first entry)
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Matches 13; Conservative
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(LIUW/) LIU W.
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                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotides comprising a contiguous sequence of 21 bp fully defined in the specification (ADPB3646) or a sequence which has one-base mismatch with ADPB3646, where the ribose residue of at least one nucleotide is protected at the 2'-0- position by 2, 4-dinitrophenyl (DNP) and where the oligoribonucleotide is capable of down-regulating the expression of the invention has expuesive the invention has cytostatic activity, and may have a use in antisense therapy. The oligoribonucleotides are useful for down-regulating the expression of the RI-alpha subunit of protein kinase A, reducing growth of cells expression of the Allpha/PKA, detecting overexpression of the RI-alpha subunit of cancer cells. The present sequence represents an oligoribonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Poly-DNP-RNA-21 5-base mismatched strand oligoribonucleotide SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            вв; oligoribonucleotide; RI-alpha subunit; protein kinase A; cytostatic;
Gaps
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3; Indels
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0; Mismatches
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                                                               1007 TGCACAAGATCTCCCGCTTC 1026
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17; Conservative
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ADP48126 ID ADP4 XX AC ADP4 XX

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This invention relates to a novel isolated polynucleotide comprising a nucleic acid sequence, the human MRCKI gene located at position 11q13, and the novel human protein kinase MRCKI encoded by it. The sequence of the invention has sequence homology to rat myotonic dystrophy kinase-related Cdc42 binding kinase (MRCK). The invention may be useful for disgnosing, prognosing, and treating kinase-related diseases, preferably diseases associated with aberrant expression of MRCKI. The present sequence is that of a DNA sequence which is part of the human MRCKI gene which may be a target for a short inhibitory (siRNA) sequence and which is related to the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic form at directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                      New human protein kinase MRCK1 with 65% sequence homology to rat myotonic dystrophy kinase-related Cdc42-binding kinase (MRCK), useful in diagnostics and as a drug target.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 5 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
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                    07-NOV-2003; 2003WO-US035609
                                                         27-NOV-2002; 2002US-0429381P
                                                                                                                                                                                                                     WPI; 2004-461109/43.
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                                                                                                                                                                             Wu L;
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0.4%; Scorè 15.2; DB 1; Length 21; 85.0%; Pred. No. 1.6e+03; tive 0; Mismatches 3; Indels

Query Match 0.4 Best Local Similarity 85.0 Matches 17; Conservative

1354 GAGATGATGAAGATGATCGG 1373

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Sequence 21 BP; 5 A; 3 C; 8 G; 5 T; 0 U; 0 Other;

This invention relates to a novel isolated polymucleotide comprising a nucleic acid sequence, the human MRCKI gene located at position 11q13, and the novel human protein kinase MRCKI gene encoded by it. The sequence of the invention has sequence homology to rat myotonic dystrophy kinase-related Cdc42 binding kinase (MRCK). The invention may be useful for disgnosing, prognosing, and treating kinase-related diseases, preferably diseases associated with aberrant expression of MRCKI. The present sequence is that of a DNA sequence which is part of the human MRCKI gene which may be a target for a short inhibitory (siRNA) sequence and which is related to the invention. Note: The sequence data for this patent did not form part of the printed appecification but was obtained in electronic form at directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

New human protein kinase MRCK1 with 65% sequence homology to rat myotonic dystrophy kinase-related Cdc42-binding kinase (MRCK), useful in diagnostics and as a drug target.

SEQ ID NO 154; 92pp; English.

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The invention describes a protein variant of the Yin Yang-1 transcription detector (1), having a 411 amino acid (aa) sequence (4) reproduced. Also described are: protein (1a) that is a homologue of (4) and includes Arg a position 303 and Lys at position 311; peptide (11) that is a fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New variant of the Yin Yang-1 transcription factor, useful for treating e.g. diabetes and autoimmune disease, also for diagnosing predisposition and in screening for therapeutic agents.
                                                                                                                                                                                                       antidiabetic, immunosuppressive, cytostatic; Yin Yang-1, transcription factor; type 1 diabetes; transgenic; diabetes; multifunctional transcription factor; type 2 diabetes; autoimmune disease; cancer; mineral metabolism disorder; lipid metabolism disorder; rat; YY-1; PCR; primer; ss.
                                                                                                                                                                            Yin yang-1 (YY-1) associated primer #28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure, Fig 11; 193pp; German.
1 GAGATCTTGAAGGTGATCGG 20
                                                                               ADQ58920 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                       19-DEC-2003; 2003WO-EP014762.
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                                                                                                                                                                                                                                                                                                        Rattus norvegicus.
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                                                                                                                                             23-SEP-2004
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                                                                                                               ADQ58920;
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                                                                ADQ58920
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protein kinase, MRCK1; myotonic dystrophy kinase, Cdc42 binding kinase, MRCK; kinase-related disease; short inhibitory RNA; siRNA; ds; human.

07-NOV-2003; 2003WO-US035609 27-NOV-2002; 2002US-0429381P

WO2004050831-A2

17-JUN-2004.

Ното варіепв

WPI; 2004-461109/43.

Liu W,

(LIUW/) LIU W. (WULL/) WU L.

Human MRCK1 siRNA target DNA sequence SeqID154.

(first entry)

09-SEP-2004

ADP48119;

ADP48119 standard; DNA; 21 BP

1457

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(II) or (Ia) and includes the positions 303 and 311 of (4); nucleic acid (III) that encodes (I), (Ia) or (II); an antibody (Ab) directed against (II) or (Ia); methods for determining a tendency to develop type 1 diabetes; transgenic non-human mammal (A) in which the germ and somatic cells contain a nucleic acid (or segment) encoding a 411 as sequence (2), or sequences with at least 95, best 99% homology, where the homologue includes 303Met and 311Arg; best 99% homology, where the homologue includes 303Met and 311Arg; best 99% homology, where the homologue includes 303Met and 311Arg; homologues of the viral methods are useful for that are protective against diabetes. The methods are useful for treatment of factor. (I), or its homologues and peptides, also nucleic acids encoding them and antisense oligonucleotides, are useful for treatment of type 1 and 2 diabetes, autoimmune diseases, cancer and disorders of mineral and lipid metabolism. Detecting mutations in the human analogue of (4) is used to determine a predisposition for these diseases. Transgenic animals that contain the sequence encoding (4), or its homologues, are used to screen for agents protective against diabetes. This sequence represents a primer associated with the isolation and analysis of yin yang-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes a protein variant of the Yin Yang-1 transcription factor (1), having a 411 amino acid (aa) sequence (4) reproduced. Also described are: protein (1a) that is a homologue of (4) and includes Arg a position 303 and Lys at position 311; peptide (II) that is a fragment of (1) or (1a) and includes the positions 303 and 311 of (4); nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variant of the Yin Yang-1 transcription factor, useful for treating . diabetes and autoimmune disease, also for diagnosing predisposition in screening for therapeutic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antidiabetic; immunosuppressive; cytostatic; Yin Yang-1; transcription factor; type 1 diabetes; transgenic; diabetes; multifunctional transcription factor; type 2 diabetes; autoimmune disease; cancer; mineral metabolism disorder; lipid metabolism disorder; rat; YY-1; PCR; primer; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 15.2; DB 1; Length 21;
85.0%; Pred. No. 1.6e+03;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 8 A; 4 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yin yang-1 (YY-1) associated primer #140.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 11; 193pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           172 GATGACGAAGACGGGGGGGA 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GACGACGAGGACGAGGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADOS9032 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYGR ) UNIV GREIFSWALD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rattus norvegicus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        402004056857-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kloeting I,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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(III) that encodes (1), (Ia) or (II); an antibody (Ab) directed against (1) or (Ia); methods for determining a tendency to develop type 1 diabetes; transgenic non-human mammal (A) in which the germ and somatic calls contain a nucleic acid (0 to segment) encoding a 411 as sequence (2), or sequences with at least 95, best 99% homology, where the homologue includes 303Met and 311Arg; and use of (A) to screen for compounds (B) that are protective against diabetes. The methods are useful for modulating activity of the YY1 (Yin Yang-1) multifunctional transcription factor. (I), or its homologues and peptides, also nucleic acids encoding them and antisense oligonucleotides, are useful for treatment of type 1 and 2 diabetes, autoimmune diseases, cancer and disorders of mineral and lipid metabolism. Detecting mutations in the human analogue of (4) is used to determine a predisposition for these diseases. Transgenic animals that contain the sequence encoding (4), or its homologues, are used to screen for agents protective against diabetes. This sequence represents a primer associated with the isolation and analysis of yin yang-1
                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mononucleotide repeat locus; human; BAT25; probe; microsatellite; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for analysing a target nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15.2; DB 1; Length 21; 35.0%; Pred, No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "Labelled with Fluorescein"
                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 1 A; 8 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mononucleotide repeat locus BAT25 probe #1.
                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                      172 GATGACGAAGACGGGGAGGA 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21 GACGACGAGGACGACGAGGA 2
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Les 17; Conservative
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                                                                                                                                                                                                                                                                                             transcription factor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hereditary cancer.
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ABN83378/
     02000000000000000x0
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comprises hybridising a polynucleotide probe comprising a segment complementary to a non-repetitive region and a segment complementary to an adjacent repetitive region, where the second segment consists of a defined number of repeats, and determining the melting point temperature of the hybrid. The method is used to analyse microsatellites, especially hereditary tumours. Alternatively, the method is used to identify an individual in a population. The present sequence is a probe for Mononucleotide repeat locus BAT25, and was used to illustrate the
          The method
                                                                                                                                                                                                                                                                                        Sequence 29 BP; 26 A; 2 C; 0 G; 1 T; 0 U; 0 Other;
consisting of repetitive and
                                                                                                                                                                                                                                                invention
                                                                                                                                                                                                                                                                                                                                           Query Match
     acid
  8X8888888888X8
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Gaps . ch 0.4%; Score 15.2; DB 1; Length 29; 1 Similarity 71.4%; Pred. No. 2.1e+03; 20; Conservative 0; Mismatches 8; Indels 3308 GATTITICITIAGGAGATTIAITITIG 3335 Local Similarity Matches ð

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28 GATTITITITITITITITITITITITITIG 1 g

RESULT 1461

AAF74918 standard; DNA; 29 AAF74918/

BP.

AAF74918

(first entry) 23-MAY-2001

tract sequence SEQ ID NO:15. CD40L poly-A Human, CD40L, promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.

Homo

WO200119844-A1.

22-MAR-2001.

13-SEP-2000; 2000WO-US024966.

99US-0153625P. 13-SEP-1999; (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.

Li Y; Crow MK, WPI; 2001-244776/25.

New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.

Example 1; Fig 3; 90pp; English.

The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (1) for CD40 ligand, having residues 313-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding antirheumatic, lamunosuppressive and antirhilammatory activities, and can be used in gene therapy. (1) is useful in the study, diagnosts and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention

Sequence 29 BP; 23 A; 3 C; 0 G; 3 T; 0 U; 0 Other;

AAF74935 standard; DNA; 29

AAF74935 RESULT 1463 AAF74935/C ID AAF74935 XX AC AAF74935

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF7405 where A in the wild type sequence at position 331 (corresponding antirheumatic, immunosuppressive and antirheumatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treatment of inflammatory and autoimmune diseases, as well as diseases ir which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                  Human, CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.
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   Score 15.2; DB 1; Length 29;
Pred. No. 2.1e+03;
0; Mismatches 8; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 29 BP; 23 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED
                                                                3258 AAGATATTTTATTTGCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3258 AAGATATTTATTTGCCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                      CD40L poly-A tract sequence SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAGGTTTTTTTTTTTTTTTTTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Fig 3; 90pp; English.
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0
                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-SEP-2000; 2000WO-US024966.
    0.4%;
Query Match 0.4%;
Best Local Similarity 71.4%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.48;
                                                                                                                                                                         AAF74907 standard; DNA; 29
                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-244776/25.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Li Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                             28
                                                                                                                                                                                                        AAF74907;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Crow MK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28
                                                                                                                                          RESULT 1462
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The present invention describes an isolated, purified nucleic acid, which residues 311-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, antirhematic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD401, is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD401 poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                          New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.
                                                                                     Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.2; DB 1; Length 29;
Pred. No. 2.1e+03;
0; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 29 BP; 23 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                 (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.
                                                     CD40L poly-A tract sequence SEQ ID NO:32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ch 0.4%;
1 Similarity 71.4%;
20; Conservative (
                                                                                                                                                                                                                                                                           13-SEP-2000; 2000WO-US024966
                                                                                                                                                                                                                                                                                                               99US-0153625P
                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-244776/25.
                                                                                                                                                                                                                                                                                                                                                                                      Crow MK, Li Y;
                                                                                                                                                                                                    WO200119844-A1.
                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                               13-SEP-1999;
               23-MAY-2001
                                                                                                                                                                                                                                      22-MAR-2001
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The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (1) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF74965 where A in the wild type sequence at position 331 (corresponding to position -1255) is replaced with C. (1) has antiarthritic, antirhematic, immunosuppressive and antinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention

New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.

Example 1; Fig 3; 90pp; English.

(NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.

WPI; 2001-244776/25.

Li Y;

Crow MK,

99US-0153625P

13-SEP-1999;

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Gaps

. 0

0.4%; Score 15.2; DB 1; Length 29; 71.4%; Pred. No. 2.1e+03; tive 0; Mismatches 9; Indels

3258 AAGATATTTTATTTGCCTTTGTCCTTTTT 3285

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Gaps

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3258 AAGATATTTTATTTGCTTTGTCCTTTTT 3285

Best Local Similarity

Matches

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28 AAGGTTTTTTTTTTTTTTTTTTT

AAF74921 standard; DNA; 29 BP

RESULT 1464

Matches 20; Conservative

Best Local Similarity

Query Match

Sequence 29 BP; 23 A; 3 C; 0 G; 3 T; 0 U; 0 Other;

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New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                                                                                                                                         Human, CD40L, promoter, CD40 ligand promoter, rheumatoid arthrividagnosis, antiarthritic, antirheumatic, immunosuppressive, antiinflammatory, inflammatory disease, autoimmune disease, ds.
                                                                                                                                                                                                                                                                                                                                   (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.
                                                                                                                                   CD40L poly-A tract sequence SEQ ID NO:25
28 AAGGITTTTTTTTTTTTTTTTTTTT 1
                                                                                                                                                                                                                                                                                    13-SEP-2000; 2000WO-US024966.
                                                                                                                                                                                                                                                                                                           99US-0153625P
                                                            AAF74928 standard; DNA; 29
                                                                                                           23-MAY-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-244776/25.
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                                                                                                                                                                                                                                                                                                                                                            Crow MK,
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Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.

WO200119844-A1

22-MAR-2001

Homo sapiens

CD40L poly-A tract sequence SEQ ID NO:18.

23-MAY-2001 (first entry)

AAF74921;

1 label, immobilising ON and PN hybrids (modified with 2 different labels nor 1 label) to the solid phase and detecting immobilised ON or PN, or the non-immobilised labelled product, directly or via an enzyme labelled antibody. The method can be used to detect specific mRNA in human, animal or plant samples. The method is easy to perform and suitable for quantitation, does not require preliminary RNA isolation, can be made quantitative and has a low background. (Updated on 25-MAR-2003 to correct PR field.)

8888888888888

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Gaps

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Indels

Length 30;

Score 15.2; DB 1; Pred. No. 2.2e+03;

Query Match 0.4%; Best Local Similarity 71.4%; Matches 20; Conservative (

0; Mismatches

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Sequence 30 BP; 2 A; 3 C; 4 G; 21 T; 0 U; 0 Other;

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The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (I) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 311 (corresponding to position -125) is replaced with C. (I) has antiarthritic, antirheumatic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer; CAT RNA; detection; quantification; determination; ss.
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                                                                                                                                                                                                                        Score 15.2; DB 1; Length 29; Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Downstream primer for synthetic full length CAT sense RNA
                                                                                                                                                                                                                                                   8; Indels
                                                                                                                                                                                               Sequence 29 BP; 23 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fritton H;
                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                          3258 AAGATATTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                          AAGGTTTTTTGTTTTTTTTTTTTTTTT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BOEF ) BOEHRINGER MANNHEIM GMBH.
            Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                 AAT69677 standard; DNA; 30 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96EP-00120480.
                                                                                                                                                                                                                      Query Match 0.4%;
Best Local Similarity 71.4%;
Matches. 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hinzpeter M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                 19-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Leying H,
                                                                                                                                                                                                                                                                                                   28
                                                                                                                                                                                                                                                                                                                                                                                            AAT69677;
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Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.

CD40L poly-A tract sequence SEQ ID NO:5.

(first entry)

23-MAY-2001 AAF74908;

AAF74908 standard; DNA; 30

(NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.

Li Y;

Crow MK,

99US-0153625P

13-SEP-1999;

13-SEP-2000; 2000WO-US024966.

WO200119844-A1

Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 24 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rheumatoid arthritis.
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Matches 20; Conserv
AAF74908/
ID AAF74908/
AC AAF7
XX AAF7
XX AAF7
XX AAF7
XX Huma
XX Huma
XX Homo
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The present sequence is a PCR primer for synthetic full length CAT sense determination of specific oligonucleotides (ON) or polynucleotides (PN). The method comprises combining a sample containing RNA or single stranded DNA with a lysis/hybridisation buffer, treating the homogenised solution with 1 or more probes, which are essentially complementary to the ON or PN and contain at least 2 different labels, one a specific binding group and the other a detectable chemical group, hybridising under stringent conditions, diluting the mixture and adding an agent that cleaves single stranded nucleic acid into monucleotides or ON, transferring the mixture to a vessel or well, coated with a substance that binds at least

Detection and quantitation of nucleic acid using probe with label and binding group - and after hybridisation treatment with RNase and capture on solid phase, particularly for RNA.

WPI; 1997-322152/30.

Example 2; Page 7; 13pp; German.

AAQ43973;

RESULT 1468 AAQ4397

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isolating a nucleic acid having a homopolyments sequence or repetitive element or conserved nucleotide sequence. (M1) comprises treating a semple containing nucleic acid compounds with an locked nucleic acid apply of the containing an uncleic acid compounds with an locked nucleic acid acid having the homopolyments sequence or repetitive element or conserved nucleotide sequence. (M1) is useful for detecting and isolating nucleic acids released from a lysed complex bological mixture comprising nucleic acids. The present sequence is a LNA capture probe, used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting and/or isolating nucleic acid molecule having homopolymeric sequence or repetitive element or conserved nucleotide sequence involves treating sample containing nucleic acid compounds with locked nucleic
                                                                                                                                                  /mod_base= OTHER
/note= "10-mer deoxy-thymine and 5-mer non-base (t10-
NBS)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method (M1) for detecting and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human reproductive system related antigen DNA SEQ ID NO: 10176.
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                   88
                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "5' AQ2, where AQ is anthraquinone"
16. 35
/*tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                 Detection; isolation; locked nucleic acid; LNA; probe;
                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "Optionally LNA nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 35 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
0.4%; Score 15.2; D
Best Local Similarity 68.0%; Pred. No. 2.4e
Matches 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3310 TITITCTTTAGGAGATTTATTTT 3334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 TTTTTTTTTNNNNNTTTTTTTT 25
                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 23; Page 67; 104pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JUN-2002; 2002US-0390928P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUN-2003; 2003WO-IB006354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL07488 standard; DNA; 38
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                                                                                                                       *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acid oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EXIQ-) EXIQON AS
                                                                                                                                                                                                                                                                                                                                                                                                           WO2004020575-A2
                                                                                                 Key
modified_base
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                                                                                                                                                                                                                           modified base
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                                                           Synthetic
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ID AAL07488/C
XC AAL074
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DT 21-NOV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence is that of an oligonucleotide, I, which is able to form a triple helix with a duplex nucleic acid (dsNA) contg. a target sequence which comprises at least one pyrimidine tract, and at least one adjacent purine tract. It is useful for therapeutic or diagnostic control of gene expression, e.g. suppression of mNA synthesis from a target gene. A specified application is targetting of RNA in the HIV-1 genome. When appropriately labelled it may also be used as a probe. Attachment of
                                                                                                                                                                                                                                                                                                                                                Purine; pyrimidine; tracts; intramolecular triplex; therapeutic; diagnostic; control; gene expression; mRNA synthesis suppression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligo:nucleotide(8) forming triple helix with target nucleic acid contain purine and pyrimidine tracts in specific orientations, useful therapeutically or diagnostically e.g. for inactivating HIV RNA, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cleavage agents caused permanent inactivation of the target by sitespecific cleavage. (Updated on 25-MAR-2003 to correct PN field.)
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                     3285
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                                                           29 AAGGTTTTTTTTTTTTTTTTTT 2
                                                                                                                                                                                                                                                                                                         Triple helix forming oligonucleotide I.
                   3258 AAGATATTTATTTGTCCTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 47; 101pp; English
                                                                                                                                                               AAQ43973 standard; DNA; 32 BP.
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92US-00826934
                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                            (revised)
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21-JAN-1992;
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28-OCT-1993
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Gaps

ADL33740;

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RESULT 1469 ADL33740

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25-SEP-2000; 2000US-0234997P.
26-SEP-2000; 2000US-0234998P.
27-SEP-2000; 2000US-0235844P.
27-SEP-2000; 2000US-0235834P.
29-SEP-2000; 2000US-0235834P.
29-SEP-2000; 2000US-0235834P.
29-SEP-2000; 2000US-0235834P.
29-SEP-2000; 2000US-0236327P.
29-SEP-2000; 2000US-0236327P.
29-SEP-2000; 2000US-0236327P.
29-SEP-2000; 2000US-0236327P.
29-SEP-2000; 2000US-0236328P.
29-SEP-2000; 2000US-0236328P.
29-SEP-2000; 2000US-0236328P.
29-SEP-2000; 2000US-0236368P.
29-SEP-2000; 2000US-0236368P.
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29-SEP-2000; 2000US-0236368P.
29-SEP-2000; 2000US-0236368P.
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20-CCT-2000; 2000US-0236368P.
20-CCT-2000; 2000US-0236368P.
20-CCT-2000; 2000US-0236369P.
20-CCT-2000; 2000US-0236369P.
20-CCT-2000; 2000US-0236613P.
20-CCT-2000; 2000US-02346474P.
20-CCT-2000; 2000US-02346474P.
20-CCT-2000; 2000US-02346474P.
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20-CCT-2000; 2000US-02346474P.
20-CCT-2000; 2000US-0234632P.
20-CCT-2000; 2000US-02346474P.
20-CCT-2000; 2000US-02346478P.
20-CCT-2000; 2000US-02346523P.
20-CCT-2000; 2000US-0234923P.
20-CCT-2000; 2000US-0234923P.
20-CCT-2000; 2000US-0234923P.
20-CCT-2000; 2000US-0234923P.
20-CCT-2000; 2000US-0234923P.
20-CCT-2000; 2000US-0234929P.
20-CCT-2000; 2000US-0234929P.
20-CCT-2000; 2000US-0234929P.
20-CCT-2000; 2000US-023593P.
20-CCT-2000; 2000US-023593P.
20-CCT-2000; 2000US-023593P.
20-CCT-2000; 2000US-023593P.
20-CCT-2000; 2000US-0234929P.
20-CCT-2000; 2000US
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2000US-0251868P.
2000US-0251869P.
2000US-0251899P.
2000US-0251990P.
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08-DEC-2000;
08-DEC-2000;
11-DEC-2000;
              Human; reproductive system related antigen; reproductive system disorder; cancer; gene therapy; ds.
                                                                                                                                                                                                                             31-JAN-2000; 2000US-0119065P.
24 FEB-2000; 2000US-0118664P.
02-MRR-2000; 2000US-0118650P.
16-MAR-2000; 2000US-0118630P.
16-MAR-2000; 2000US-01189174P.
17-MAR-2000; 2000US-01190076P.
18-MAR-2000; 2000US-01190076P.
19-MAY-2000; 2000US-01191076P.
19-MAY-2000; 2000US-01191076P.
19-MAY-2000; 2000US-01191076P.
19-MAY-2000; 2000US-0119115P.
07-JUL-2000; 2000US-021664P.
11-JUL-2000; 2000US-021664P.
11-JUL-2000; 2000US-021664P.
11-JUL-2000; 2000US-021890P.
11-JUL-2000; 2000US-021890P.
11-JUL-2000; 2000US-021890P.
14-AUG-2000; 2000US-022514P.
16-AUG-2000; 2000US-022314P.
16-AUG-2000; 2000US-022314P.
16-AUG-2000; 2000US-022314P.
16-AUG-2000; 2000US-022314P.
16-AUG-2000; 2000US-023314P.
                                                                                                                                                                                         17-JAN-2001; 2001WO-US001339
                                                                                                             WO200155320-A2
                                                                       Homo sapiens
                                                                                                                                                      02-AUG-2001
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Gaps

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Rosen CA,

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method of the invention has cytostatic and virucide applications and may be useful for treating HPV infections or HPV-associated conditions including cervical dysplasia, cervical carcinoma, oral mucosal cancer and laryngeal cancer. Furthermore, the method may be utilised during vaccine production and gene therapy. The current sequence is that of the HPV-targeted ribozyme (Rz)-related RNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Compsn. contg. 5-thiolated (Oligo-poly-)-nucleotide(s) - for treating HIV infection, AIDS and for preventing HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The oligomer comprises a non-thiolated (binding) homooligonucleotide region ( d(A)12 ) to promote the binding of the remaining portion of the S-thiolated oligonucleotide ( MdU 28 ) to a homopurine site of the viral genome via triple-helix formations. The oligo is used to in the treatment of HIV. See also AAQ22624-27 and AAQ25017-Q25031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV; thiolation; reverse transcriptase; primer; inhibition; homooligomer;
                                                                                                                                                                                                           0.4%; Score 15.2; DB 1; Length 39; 63.9%; Pred. No. 2.5e+03; tive 0; Mismatches 13; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                h 5:4%; Score 15.2; DB 1; Length 40; Similarity 14.3%; Pred. No. 2.6e+03; 4; Conservative 16; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 40 BP; 12 A; 0 C; 0 G; 0 T; 28 U; 0 Other;
                                                                                                                                                                                                                                                                                                          3308 GATITITICITIAGGAGATITATITITIGGACTICAA 3343
                                                                                                                                                                    Sequence 39 BP; 32 A; 2 C; 2 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                        39 GACTITITITITITITITITITITITITITITITICAA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide specific for HIV proviral DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3258 AAGATATTTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schinazi RF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAAAAAUUUUUUUUUUUUUUUUUUUUUUUU 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 11; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ33681 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aradi J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              91WO-US005919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90US-00568131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ25032 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYNY-) RES FOUND UNIV NEW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-JUL-1992 (first entry)
                                                                                                                                                                                                                                                              23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-096567/12.
                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ho YK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9203127-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bardos TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loc
Matches
                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel method for treating Human papillomavirus (HPV) infection comprising administering to a patient a nucleic acid molecule that inhibits expression associated with HPV replication. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HPV infection, replication, cytostatic, virucide, cervical dysplasia,
carcinoma, oral mucosal cancer, laryngeal, vaccine, gene therapy,
double internal trans-acting ribozyme, single, dITRz, ITRz, CLIP portion,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3' flanking RNA of IT ribozyme (Rz) from CLIP portion of SNIPAA cassette.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating human papilloma virus (HPV) infection, e.g. cervical dysplasias or HPV-associated cervical carcinoma, comprises administering to a patient a nucleic acid molecule that inhibits expression associated with HPV replication.
                                                                                                                                                                                             antigen is
                                                                                                                                                                                                                                                                                                          The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                          Isolated nucleic acid molecule encoding a reproductive system anused in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 10176; 1297pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 0.4%; Score 15.2; DB 1; Length 38; 1 Similarity 63.9%; Pred. No. 2.5e+03; 23; Conservative 0; Mismatches 13; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3299 TITCIAIAGGAITITITITITAGGAGAITITATITITI 3334
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 38 BP; 35 A; 0 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thiboutot D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rriciricerririririririr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 54; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Christensen N,
                                                                                                   Ruben SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-JUN-2002; 2002US-0391795P:
14-OCT-2002; 2002US-0417997P.
21-FEB-2003; 2003US-0449066P.
         05-JAN-2001; 2001US-0259678P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JUN-2003; 2003WO-US020340.
                                                     (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PENN-) PENN STATE RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADI32814 standard; RNA; 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                        protein of the invention
                                                                                                   Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pan W,
                                                                                                                                               WPI; 2001-465570/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNIPAA cassette; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-082869/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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22-APR-2004

ADI32814;

RESULT 1471 ADI32814/c

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Query Match

Best Loc Matches

Unidentified

08-JAN-2004

Clawson GA,

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Gaps

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The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine Mbol DNA fragments of between 250 and 500 by with an (AClis and a (TC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and Mbol sites, the frequency of (TG)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of proponentially immortant traits and in cartie to allow selective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             economically important traits esp. In cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic oligonucleotide, dinucleotide repeat, cytostatic, apoptosis, cell cycle arrest, cell proliferation, caspase, cytokine, interleukin, tumour necrosis factor; TNF; cancer, carcinoma; sarcoma; leukemia;
                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic bovine DNA markers - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match 0.4%; Score 15; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 1.2e+03; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 0 A; 0 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Table 7; Page 235; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                  mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2319 GTGTGTGTGTGTG 2333
                                                                                                                                               92WO-US000340
                                                                                                                                                                                            91US-00642342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-DEC-2000; 2000WO-CA001467.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic oligonucleotide 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0170325P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH46010 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                              Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                            WPI; 1992-284684/34.
                                                                                                                                                                                                                                               (GENM-) GENMARK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200144465-A2.
                                                  WO9213102-A1.
                                                                                                                                            15-JAN-1992;
                                                                                                                                                                                            15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .ymphoma; ss
                                                                                               06-AUG-1992
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     Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH46010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1475
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MboI DNA fragments of between 250 and 500 by with an (ACI)5 and a (TCI)5 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellite is summarised in the sequence information of the microsatellite sequence were used to generate the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence were used to generate the microsatellite (using the program OFTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                   PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 15.2; DB 1; Length 41;
85.0%; Pred. No. 2.6e+03;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 41 BP; 6 A; 0 C; 15 G; 20 T; 0 U; 0 Other;
                                                                                                                   Microsatellite sequence from clone TGLA122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Microsatellite sequence from clone TGLA171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2824 ATATATACATATATATAT 2843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Table 7; Page 202; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 ACACACACATATATATAT 1
                                                                                                                                                                                                                                                                                                                                                                                      92WO-US000340.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      91US-00642342.
                                          (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-284684/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                         15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-1991;
                                          25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                         WO9213102-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                                                                         06-AUG-1992
                                                                                                                                                                                                                                            30s taurus.
AAQ33681;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ33764;
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Matches

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Composition comprising synthetic oligonucleotides which comprise multiple repeats of dinucleotides such as GT, TG useful for treating cancer by inducing cell cycle arrest, inhibiting proliferation, activating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     marker gene; tumour; Kaposi's Sarcoma; peripheral blood mononuclear cell; PBMC; expressed keratin 14; TIE 1; Salioadhesin; Siglec 1; angiogenesis;
                                                                                                                                                                                                                                                 The present sequence is that of a synthetic oligonucleotide useful to the invention. The invention relates to a composition, comprising a 2 to 20 base 3-OH, 5'-OH synthetic oligonucleotide which comprises multiple repeats of dinucleotides such as GT. TG, etc., according to specific formula and having cytostatic activity. The oligonucleotide compositions are useful for inducing cell cycle arrest, inhibition of proliferation, activation of apoptosis or production of cytokines such as interleukin (IL)-1-beta, IL-6, IL-10, IL-12 and tumour necrosis factor (TNF)-alpha by immune system cells, in an animal having cancer such as primary carcinoma, secondary carcinoma, primary sarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                      and secondary sarcoma such as, leukemia, lymphoma, breast, prostate, colorectal, ovarian or bone cancer. The compositions induce apoptosis independent of Fas, p53/p21, p21/waf-1/CIP, p15(ink4B), p16(ink4), drug resistance, caspase 3, transforming growth factor (TGF)-beta 1 receptor and hormone dependence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determining whether a treatment is effective in changing a status of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KS3 and KS4 SAGE library over-expression showing tag, SEQ ID No 10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15; DB 1; Length 15;
Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 15 BP; 0 A; 0 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      drug target; tag; SAGE library; KS3; KS4; BB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Pred. No. 1...
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                                                                                                                                                                                                                    Claim 5; Page 17; 77pp; English
                               (BION-) BIONICHE LIFE SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2319 GTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC13343 standard; DNA; 15 BP
29-AUG-2000; 2000US-0228925P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-SEP-2001; 2001EP-00203703.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PRIM-) PRIMAGEN HOLDING BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                 Phillips NC, Filion MC;
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                                                                                                  WPI; 2001-398150/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van Der Kuyl AC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-DEC-2003
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                                                                                                                   The invention relates to a novel method for determining whether a treatment is effective in changing a status of a certain set of target cells in an individual. The method comprises obtaining a sample from an individual after intilation of the treatment; and determining whether the sample comprises an expression product of at least one marker gene. The marker gene and a proteinaceous molecule (which can bind to the protein derived from the marker gene of the invention) are useful for determining whether a treatment is effective in counteracting a tumour in an individual, especially Kaposi's Sarcoma. Peripheral blood mononuclear cell (PBMC) expressed keratin 14, TIE 1, Salioadhesin, or Sighec 1 sequences or a fully defined sequence given in the specification, or
certain set of target cells in an individual comprises determining whether the sample comprises an expression product of at least one marker
                                                                                                                                                                                                                                                                                                                                                their analogues are useful as indicators for angiogenesis and for detecting the presence of a tumour cell in an individual. The expression product of a gene comprising a marker gene of the invention is useful as a drug target. The compound is useful for preparing a medicament. This polymucleotide sequence represents a tag sequence which showed overexpression in Kaposi's Sarcoma SAGE libraries KS3 and KS4 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; T-cell associated disease; Wbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; Apperensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; disease; presense atrophic gastritis; disease; Apperensitivity disease; multiple sclerosis; Alzehiner, disease; Apperensitivity; doodpasture's syndrome; type II hypersensitivity; doodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; tilly; fungal infection; Candida; parasitic infection; echistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15; DB 1; Length 15;
100.0%; Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 3 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Prec. ...
                                                                                   Disclosure; SEQ ID NO 10; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human Vbeta gene repeat sequence #139.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3704 CATGGTGGCCAGAGG 3718
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95US-00531241.
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2002150891-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nvention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADH70349;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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including autoimmune diseases, degenerative nervous system diseases, graft versus host diseases, hypersensitivity diseases, infectious diseases, and neoplastic diseases. Autoimmune diseases include Addison's disease, atrophic gastritis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include Type allergies, Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections manifested in leprosy. Infectious diseases include viral infections the yeast genus Candida, parastitic infections such as those caused by schistcosmes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases
                                                                                    Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity; disease; type I hypersensitivity; doodpasture's syndrome; type II hypersensitivity; doodpasture's syndrome; type II hypersensitivity; leprosy; infectious disease; viral infection; HIV; fungal infection, Candida; parasitic infection; shibstoosme; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                           The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, VbetaRNA or cDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.2e+03; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 7 A; 0 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                   Disclosure; SEQ ID NO 543; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Vbeta gene repeat sequence #141.
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Best Local Similarity 100.C
Matches 15; Conservative
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                                          WPI; 2004-059052/06.
Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         breast cancer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JS2002150891-A1
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                                                                                                                                                        Vbeta gene
Hood LE,
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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, VbetaRNA or cDNA. The kit is useful for diagnosing organ transplant compressing and treating T-cell associated diseases including autoimmune disease, degenerative nervous system diseases. Complexit diseases, hypersensitivity diseases, infectious diseases and neoplastic diseases. Hypersensitivity diseases include Multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple collections and Alzheimer and contact with allergens that lead to allergies, Type II hypersensitivities such as those contact with allergies und as those caused by viruses such as HIV, fungal infections such as those caused by viruses such as HIV, fungal infections such as those caused by chieves and bacterial infections such as those caused by chieves tillaria and bacterial infections such as those caused by chieves tillaria and bacterial infections such as those caused by Nycobacterium. Neoplastic diseases include as concer of the brain, however the allergens include as concer of the brain, however the allergens and cancers such as concered the brain, however the such as concerned the present of the brain, however the past genus caused by the contact of the brain, however the such as concerned the present of the brain, however the past genus caused by the contact of the brain, however the present of the brain, however the such as concerned the present of the brain, however the present of the present 
                                                                                                                                                                                                                                                                             Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eating disorder, polymorphism, dataset, allele, HGBASE identification, serotonin receptor 1D; delta-opioid receptor; dopamine receptor D2; anorexia nervosa; bulimia nervosa; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           breast. The present sequence represents a Vbeta gene repeat sequence.
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Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 545; 164pp; English.
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100.0%; Pred
0; N
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                     94US-00309335.
95US-00531241.
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Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                               WPI; 2004-059052/06
                                                                                                                                                                            Hood LE, Rowen L;
                                                                                              (ноор/) ноор г Е.
                                                                                                                         (ROWE/) ROWEN L.
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                     19-SEP-1994;
19-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                              Vbeta gene
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%XXCCCCCCCCCCCCCCCXX8X11111XXBXXBXXBXXBXXBXX
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16-JUL-2001; 2001US-0305153P.

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Example 1; Page 77; 173pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200268637-A2.
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                                                                                                                                                                                                The invention relates to a novel isolated nucleic acid molecule comprising a variant gene associated with an eating disorder and selected from any of 119 polymorphisms with their corresponding genotyping in dataset, alleles and HGBASE identification, given in the specification. The novel nucleic acid molecule has polymorphisms in the serotonin receptor 1D, delta-opioid receptor, or dopamine receptor D2, which is useful in diagnostic and prognostic assays for eating disorders, in particular ancrexia nervosa and bullania nervosa. This polynucleotide sequence represents a dopamine D2 receptor probe of the invention
                                                                                                                       nucleic acid molecule having polymorphisms in the serotonin receptor delta-opioid receptor, or dopamine receptor D2, useful in diagnostic prognostic assays for eating disorders, such as anorexia and bulimia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; 5'-anchored; ss.
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                                                                                                                                                                                                                                                                                                                                . Match 0.4%; Score 15; DB 1; Length 16; Local Similarity 100.0%; Pred. No. 1.3e+03; es 15; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                              Sequence 16 BP; 2 A; 9 C; 3 G; 2 T; 0 U; 0 Other;
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20-JUL-2001; 2001US-0306440P.
13-NOV-2001; 2001US-0331285P.
19-DEC-2001; 2001US-0340843P.
19-DEC-2001; 2001US-0340844P.
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                                                        PRIC-) PRICE FOUND L'ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vogel JM;
                                                                            Bergen AW, Yeager M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-277795/28.
                                                                                                WPI; 2003-268122/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JAN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-JUN-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT27921;
                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                         nervosa
                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1480
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Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor directed primer; comprising a sequence complementary to an adaptor segment. The present sequence is an example of a SSR primer. The method represents a modified fragment length polymorphism assay, which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprofective; antibacterial; replication; pancreatitis; encephalltis; myocarditis; meningitis; infection; hepatitis; liver fallure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                  is partic. useful for genome fingerprinting, i.e. for genetic trait marking and germplasm comparisons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15; DB 1; Length 17; 00.0%; Pred. No. 1.38+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 17 BP; 7 A; 8 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WNV Amberzyme substrate SEQ ID NO 6106.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 15; Conservative
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A; 5 C; 6 G; 0 T; 2 U; 0 Other;

Seguence 17 BP; 4

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least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, waycarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least tenz 2-0-methyl modifications, phosphorothiadre linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid molecule that modulates replication of West Nile Virus
                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver fallure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                      0.4%; Score 15; DB 1; Length 17; 73.3%; Pred. No. 1.3e+03; tive 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WNV minus strand Inozyme substrate SEQ ID NO 11302.
                                                                                                                                  Seguence 17 BP; 2 A; 6 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 23; SEQ ID NO 11302; 495pp; English.
                                                                                                                                                                                                                                                  1836
                                                                                                                                                                                                                                                                                                                                                                                 BP
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|:||:||:||||||:|
CUGCUCUGGGAGAUC 15
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ACN11299 standard; RNA; 17
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                                                                                               molecule of the invention
                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-706994/76.
                                                                                                                                                                                       Similarity
11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           West Nile Virus
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                                                                                                                                                                                                                                                                                                                                                                                                                    ACN11299;
                                                                                                                                                                      Query Match
Best Local &
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(BLAT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid molecule that modulates replication of West Nile Virus
                                     Gaps
                                                                                                                                                                                                                                                                                                                                        WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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0.4%; Score 15; DB 1; Length 17;
100.0%; Pred. No. 1.3e+03;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15; DB 1; Length 17; Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                          WNV minus strand Inozyme substrate SEQ ID NO 11303.
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                                                                                                                                                                                                  ВЪ.
                                                                       1822 CTGCTCTGGGAGATC 1836
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                                                                                                                                                                                                 ACN11300 standard; RNA; 17
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                                                                                                                                                                                                                                                                      (first entry)
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                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                    Amberzyme; Zinzyme; ss
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                Local Similarity
les 15; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blatt L,
 Query Match
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Matches
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1822 CTGCTCTGGGAGATC 1836

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RESULT 1484

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proteins and their coding sequences: MD23, MD24, MD212, MD213 sencoded at chromosome 7q22.1, MD23, sencoded at chromosome 7q22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15q26.1. The MD23, MD24, MD27, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease clusted as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapoulic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                               Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to novel human zinc finger-containing
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1.3e+03;
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                                                                                                                    Human MDZ3 scanning oligonucleotide SEQ ID 667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'Match 0.4%; Score 15; DB Local Similarity 100.0%; Pred. No. 1.3 les 15; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JUL-2002; 2002EP-00016874.
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                                                                         20-NOV-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                           EP1281758-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA99677;
                           ADA99678;
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     8 X 8 X 6 X 8 X 8 X 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (NNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatis, a condition related to WNV infection e.g. pancreatis, inverted the properties, methodicis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nuclecides and a 3' end modification of a 3'-3' inverted abasic molety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                           WNV, West Nile Virus, antiinflammatory, cytostatic, hepatotropic, virucide, neuroprotective, antibacterial, replication, pancreatitis, encephalitis, myocarditis, meningitis, infection, hepatitis, liver failure, cancer, cirrhosis, Hammerhead, Inozyme, DNAzyme,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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73.3%; Pred. No. 1.3e+03;
ve 4; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 23; SEQ ID NO 6105; 495pp; English
                                                                                                                                                                                                                                                                                              WNV Amberzyme substrate SEQ ID NO 6105.
                                                                                                                                                ACN06102 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTGCTCTGGGAGATC 1836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-OCT-2001; 2001WO-US048350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-OCT-2000; 2000US-0242411P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L.
CTGCTCTGGGAGATC 2
                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 73.3
1es 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt L, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-706994/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200268637-A2.
                                                                                                                                                                                                                                                22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-SEP-2002.
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                           16
                                                                                                                                                                                                ACN06102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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ACNO 6102

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Gaps

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Indels

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Cytostatic; immunostimulant; gene therapy; vaccine; human;

ADA99678 standard; DNA; 17 BP

RESULT 1485

ઠે 요 ADA99678 ID ADA9

Human MDZ3 scanning oligonucleotide SEQ ID 666.

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New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 15; DB 1; Length 17;
100.0%; Pred. No. 1.3e+03;
ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 5 A; 5 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 665; 103pp; English.
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100.0%; F1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1529 CCGAGGAGCAGCTCA 1543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ61957 standard; RNA; 17 BP
                                      30-JUL-2002; 2002EP-00016874.
                                                                         02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAY-2002; 2002WO-US016840.
                                                                                                                                           Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cccaccaccaccrca 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 100.
1es 15; Conservative
                                                                                                                                                                                WPI; 2003-423107/40.
                                                                                                            (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200297114-A2
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     05-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-DEC-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Matches
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     The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is cancoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome second in the sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder or associated with MDZ1, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids and proteins are also useful for diagnosing or monitoring a disease acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
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zinc finger protein; MDZ3; MDZ4; MDZ1; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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100.0%; Pred. No. 1.3e+03;
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Best Local Similarity 100.0%; Pred. No. 1.3
Americative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; SEQ ID NO 666; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                02-AUG-2001; 2001US-00922181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1529 CCGAGGAGCAGCTCA 1543
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                                                                                                                                                                             30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                    Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCGAGGAGCAGCTCA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-NOV-2003 (first entry)
                                                                                                                                                                                                                                                                                                                       WPI; 2003-423107/40.
                                                                                                                                                                                                                                                 (AEOM-) AEOMICA INC
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                                                                       Homo sapiens
                                                                                                        EP1281758-A2
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                                                                                                                            The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV acid molecules are useful for reading breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ65210 - ABZ65224, ABZ65531, ABZ65210 - ABZ65224, ABZ65531, ABZ65210 - ABZ6524,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate
                                                                                                                                                                                                                                                                                                                                            Gaps
                                                         Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting, amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; 5'-anchored; ss.
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                                                                                                                                                                                                                                                                                                                 0.4%; Score 15; DB 1; Length 17; 100.0%; Pred. No. 1.3e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5'-anchored simple sequence repeat primer DBD(AC)7.5.
                                                                                                                                                                                                                                                                                             Seguence 17 BP; 2 A; 6 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          В 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 76; 173pp; English.
                                                                                                         Claim 58; Page 125; 185pp; English.
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                                                                                                                                                                                                                                                                    ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                           GGCAAGCCCCTTGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                Local Similarity 100.
Les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Morgante M, Vogel JM;
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                                 WPI; 2003-140484/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JAN-1997
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            Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
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                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                             Matches
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fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor segment. The present sequence is an example of a SSR primer, which is a flanked at its 5'-end by degenerate nucleotides. The method represents anodified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense compounds complementary to Cellular Inhibitor of Apoptosis-1 useful for e.g. diagnostics, therapeutics, and as research reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cellular Inhibitor of Apoptosis-1; antisense; diagnostic; therapeutic; c-IAP-1; prophylaxis; infection; inflammation; tumor formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human c-IAP-1 mRNA inhibiting antisense oligo ISIS #23345.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 15; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 1.4e+03; Matches 15; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 8 A; 7 C; 0 G; 0 T; 0 U; 3 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                        germplasm comparisons
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US5958772-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  apoptosis-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ22163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1490
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The invention relates to a method of inducing cellular apoptosis comprising administering to a cell an effective amount of an antisense compound targeted to a nucleic acid molecule encoding human cellular inhibitor of apoptosis-1 (cIAP-1), so that expression of cIAP-1 is inhibited and apoptosis is induced. The cell used in the method is a cancer cell. The antisense compound is an antisense oligomolecide that comprises at least one modification of the internucleoside linkage, sugar moiety or nucleobase, wherein the modification is a phosphorothioate internucleoside linkage, a 2'-O-methoxyethyl sugar moiety or a 5-methylytosine nucleobase. The antisense oligomolecitide is a chimeric oligomolectide. The methods and compositions are useful for the prevention and/or treatment of diseases or conditions associated with a pherent expression or activity of human cIAP-1, such as cancer, autoimmune disorders and viral infections. This sequence represents a human cIAP-1 DNA antisense oligomolectide of the invention.
                                                                                                                                                                                                                                                                        Human; cellular inhibitor of apoptosis-1; cIAP-1; ss; cellular apoptosis; cancer; antisense oligonucleotide; phosphorothioate internucleoside linkage; 2'-0-methoxyethyl sugar moiety; 5-methylcytosine; autoimmune disorder; viral infection; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inducing cellular apoptosis by administering an antisense modulating the human Cellular Inhibitor of Apoptosis-1, useful in preventing or treating cancer, autoimmune disorders and viral infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 4 A; 3 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                          Human cIAP-1 DNA antisense oligonucleotide #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Scor.
100.0%; Pred. No. 1...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; SEQ ID NO 21; 25pp; English.
                                                                                                                              ВР
1606 CAGAAGTGCATCCAC 1620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-DEC-1998; 98US-00205204.
16-JUN-1999; 99WO-US013624.
24-SEP-2001; 2001US-00857278.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-JUN-2003; 2003US-00464158
                                                                                                                                                                                                                                                                                                                                                   immunosuppressive; virucide
                     17 CAGAAGTGCATCCAC 3
                                                                                                                          ADM96437 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bennett CF, Ackermann EJ;
                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BENNETT C F.
ACKERMANN E J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-090476/09.
                                                                                                                                                                                                                                                                                                                                                                                                                          US2004009599-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BENN/) BENNETT
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                     17-JUN-2004
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                                                                                                                                                                 ADM96437;
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The present invention relates to a method for synthesising a branched or multiply connected macromolecular structure, comprising oligonucleotide clamps (OC). The macromolecular structure is capable of specifically binding to a target molecule, and can therefore be used as probes. At least one OC comprises a target binding sequence that binds specifically and stably with the target molecule, and at least two OCs comprise signal generation moieties capable of generating a detectable signal in the presence of the target molecule. In addition the OCs are connected to one another by thioalkylamino, or thiophosphorylalkylamino bridges. The present sequence is an OC used in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthesizing branched nucleic acids useful as diagnostic and molecular probes, involves combining first units having haloalkylamino groups and second units having thiol or phosphorothioate groups.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymphogene; Prostogene; Breastogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer; RNA interference.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human protein kinase, lysine deficient 4, SDSO target region #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15; DB 1; Length 19;
Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 9 A; 10 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Scor.
100.0%; Pred. No. 1...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 7; Col 19; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADO23027 standard; cDNA; 19 BP.
                AAF60477 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2319 GTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                            97US-00787321
                                                                                                                                                                                                                                                                                                              96US-0009918P.
                                                                                                                Oligonucleotide clamp #18
                                                                                                                                                Oligonucleotide clamp; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-201911/20.
                                                                                                                                                                                                                                                                                                                                            (FARB ) BAYER CORP.
                                                                                                                                                                             Unidentified.
                                                                                                                                                                                                                                                                                                              12-JAN-1996;
                                                                                                                                                                                                                                                                            03-JAN-1997;
                                                                                                                                                                                                               US6180777-B1
                                                                               27-APR-2001
                                                                                                                                                                                                                                               30-JAN-2001.
                                                 AAF60477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AD023027;
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AAF60477/
ID AAF6
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Gaps

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Length 18; 0; Indels

Score 15; DB 1; Le Pred. No. 1.4e+03;

1606 CAGAAGTGCATCCAC 1620

15; Conservative

Query Match Best Local Similarity Matches 15; Conservat

17 CAGAAGTGCATCCAC 3

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The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonuclectides (termed short double stranded oligonuclectides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the sequence within the abnormal gene encoding a protein or playing other composition colls and protein chip microarrays, identifying a specific DNA composition of a solution of a sequence of computer and specific Software. Composition of a sesistance of computer and specific software, composition of an RNA molecule and making sure that selected sequence of software. Also included are pharmaceutical compositions of gene drugs (such as Dermogene, Lungene, Hepatogene, Leukogene, Lumpene, Lungene, Hepatogene, Lukogene, Lumpene, Lungene, Hepatogene, Lukogene, Lumpene, Lumpene, Hepatogene, Lukogene, Lumpene, Lumpene, Lumpene, Lumpene, Lumpene, Lumpene, Lukogene, Lukokogene, Lukogene, Lukokogene, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide molecules or short interfering RNA (siRNA) (comprising identifying a special pattern that can be localised in any position of an oligonucleotide sequence evaluating the specificity of a selected sequence). The Short interfering RNA (siRNA) are targeted against genes involved in viral infection, malignant tumours, genetic and metabolic diseases. The methods are useful for designing and selecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              short double-stranded oligonucleotides as a gene drug that can specifically inactivate a group of corresponding genes. The composition
                                                                                                                                                                                                                                                                                                                                                                                                                                   Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 10; 58pp; English.
                                                                                                                                                             16-SEP-2002; 2002US-00016490.
                                                                                                                                                                                                                   16-SEP-2002; 2002US-00016490.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific DNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-355427/33.
                                                US2004072769-A1.
                                                                                                                                                                                                                                                                      YINJ/) YIN J Q.
  Homo sapiens.
                                                                                                         15-APR-2004.
                                                                                                                                                                                                                                                                                                                            Yin JQ;
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Sequence 19 BP; 1 A; 11 C; 3 G; 4 T; 0 U; 0 Other;

expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present sequence is a target region for an SDSO from an human cDNA. may be used for treating diseases or disorders associated with abnormal

ö Gaps ö 0.4%; Score 15; DB 1; Length 19; 100.0%; Pred. No. 1.5e+03; ive 0; Mismatches 0; Indels 15; Conservative Local Similarity Query Match Best Loca Matches

2589 GCTCGGCCCTCCCA 2603

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ADO22936 standard; DNA; 19 BP. (first entry) 01-JUL-2004 AD022936; RESULT 1494

Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymphogene; Prostogene; Brastogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer; Human sequence containing an SDSO cleavage site #14.

Homo sapiens.

RNA interference

JS2004072769-A1

15-APR-2004.

16-SEP-2002; 2002US-00016490.

16-SEP-2002; 2002US-00016490

(YINJ/) YIN J Q.

rin 30;

WPI; 2004-355427/33.

Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify specific DNA sequences.

Claim 12; SEQ ID NO 25; 58pp; English.

The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the gene chip and protein chip microarrays, identifying a specific DNA sequence within the abnormal gene encoding a protein or playing other biological roles with the assistance of computer and specific software, considered and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related software. Also included are pharmaceutical compositions of gene drugs (such as Dermogene, Iungene, Hepatogene, Leukogene, Lymphogene, Prostogene, Braintumogene and Skin-whitogene including but group of specific 19-25 nt daRNA, 19-25 nt daRNA and/or single-stranded RNA and/or DNA with the special pattern, 5'-cogened or practical pattern, 5'-cogened or pharmaceutical carifect condensation agents (or none), one or more nucleic acid condensation agents (or none), one or more predicting and selecting and activiting a materials and a simplified method for predicting and solditional specific and efficacious small double-stranded oligonucleotides (SDSO), antisense oligonucleotide molecules or short interfering RNA (SIRNA) (comprising identifying a special pattern that can be localised in any position of an oligonucleotide sequence evaluating the specificity of a selected sequence). The Short interfering RNA (siRNA) are targeted against genes involved in viral infection, malignant tumours, genetic and metaboolic diseases. The methods are useful for designing and selecting short double-stranded oligonucleotides as a gene drug that can Antisense compound; tumour necrosis factor receptor 1; liver disease; TNFR1; hepatitis; liver injury; hyperproliferative disorder; cancer; mouse; murine; ds.

TNFR1 expression modulation related antisense oligo SEQ ID No 176.

ABT05146 standard; DNA; 20 BP.

RESULT 14 ABT05146

11-OCT-2002 (first entry)

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ABT05146;

2 GGCATCAAGCTGCGG 16

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specifically inactivate a group of corresponding genes. The composition may be used for treating diseases or disorders associated with abnormal expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present sequence is a target region for an SDSO from an (unspecified) human gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention comprises antisense oligonucleotides that are capable of inhibiting expression of the human clusterin gene. The antisense oligonucleotides of the invention are useful for inhibiting the expression of clusterin in cells. The antisense oligonucleotides are also useful for treating an animal with a disease or condition associated with clusterin (e.g. hypercholesterolaemia; cardiovascular disorders; hypercholesterolaemia; cardiovascular disorders; hyperproliferative disorders; and hyperlipidemic disorders; The present DNA sequence represents a clusterin antisense oligonucleotide of the invention. NOTE: The present DNA sequence has a phosphorothioate backbone and also contains 2'-O-methoxyethyl wings
                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel antisense compound targeted to nucleic acid molecule encoding clusterin, useful for treating animal having disease associated with clusterin such as hyperlipidemic disorder, cardiovascular disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antisense inhibition; antisense oligonucleotide; clusterin; hypercholesteroleamia; cardiovascular disorder; ss; hyperpoliferative disorder; hyperlipidemic disorder; phosphorothioate backbone; 2'-0-methoxyethyl wing.
                                                                                                                                                                                             ö
                                                                                                                                                      0.4%; Score 15; DB 1; Length 19;
100.0%; Pred. No. 1.5e+03;
                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human clusterin inhibiting antisense oligonucleotide 34.
                                                                                                                 Sequence 19 BP; 4 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 83; 125pp; English.
                                                                                                                                                                                                                               2990 TTTCTGGCACCGCAG 3004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-SEP-2000; 2000US-00659791.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-2001; 2001WO-US028235.
                                                                                                                                                                                                                                                                                                                                                              ABN99700 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                      TTTCTGGCACCGCAG
                                                                                                                                                                                           15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Freier SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-404805/43.
                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200222635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                        16-AUG-2002
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                                                                                                                                                                                                                                                                  19
                                                                                                                                                      Query Match
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                                                                                                                                                                                           Matches
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Dean NM;

Zhang H,

Cowsert LM,

Baker BF,

WPI; 2002-583481/62.

(ISIS-) ISIS PHARM INC.

22-OCT-2001; 2001WO-US051224. 24-OCT-2000; 2000US-00695451.

WO200248168-A1. 20-JUN-2002

Mus sp.

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                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compound targeted to nucleic acid molecule encoding tumor necrosis factor receptor 1 (TNFR1), useful for treating humans having disease associated with TNFR1 e.g. hepatitis, liver injury, liver cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an antisense compound 8 to 30 nucleotides in length targeted to nucleic acid molecule encoding tumour necrosis factor receptor 1 (TNFRI), where the antisense compound inhibits expression of TNFRI. The antisense compound is useful for inhibiting the expression of TNFRI in cells or tissues. The antisense compound is also useful for treating an animal (preferably human) having a disease or condition associated with TNFRI. e.g. a liver disease (such as hepatitis, or liver injury) or a hyperproliferative disorder such as cancer, by inhibiting the expression of TNFRI. The antisense compound is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. This polynucleotide sequence represents a mouse oligonucleotide relating to the TNFRI of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 1.6e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kaposi's Sarcoma TAG019 SEQ ID NO:45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 21; Page 61; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1571 ACCAGGTGGCCCGGG 1585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABQ81895 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-NOV-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABQ81895;
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382 GGCATCAAGCTGCGG 396

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Gaps

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Query Match

Matches

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Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EPB2; LAP; TCF5; CRP2; NFL16; LIGDBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness; oxidative stress response; IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thl response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:173
                                                                                                                                                                                                                                            The invention comprises a gene screening method for groups with a high risk of developing cervical cancer. The gene screening method involves determining whether or not epithelial tissue cells have the integral human papilloma virus (HPV) 16.18 E6/F3 gene segment. The method of the invention may be used to screen high risk groups for cervical cancer. I present sequence represents a PCR primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2'
cytosines are 5-methylcytosine"
                                                                                                                                                                Gene screening method for high risk group of cervical cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15; DB 1; Length 20;
Pred. No. 1.6e+03;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 6 C; 7 G; 4 T; 0 U; 0 Other;
                                       (BEIJ-) BEIJING BOTAIDI BIOLOGICAL ENG SCI & TEC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "2'-methoxyethyl (2'-MOE) cytosines are 5-methylcytosine"
                                                                                                                                                                                                       Claim 4; Page 4 (Disclosure); 7pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Scc...
100.0%; Prec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2990 TTTCTGGCACCGCAG 3004
16-FEB-2001; 2001CN-00101870.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 rrrcreecaccecae 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL94407 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.4
Best Local Similarity 100.
Matches 15; Conservative
                                                                                  Qian D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1. .20
/*tag=
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                                                                                                                       WPI; 2002-148635/20.
                                                                                  Yan Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                              nvention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL94407;
                                                                                  Wang B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL94407
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a method for determining if an individual has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing a status of a set of target cells, comprising determining if a sample of the subject has an expression product of at least one marker gene. Also described is a compound capable of altering the expression or activity of Keratin 14, TIE 1, Salioadhesin or Siglec, and kits containing them from the present invention can be used in a diagnostic method, particularly as an indicator of angiogenesis or to determine presence of a tumour cell. The method of the invention is suitable to determine within a few days if a certain treatment against Kaposi's Sarcoma is successfull. ABQ81816 to ABQ82006 represent nucleotide sequence used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                   Determining presence of a tumor cell or angiogenesis, and the effectiveness of treatment, by detecting the presence of marker genes is useful to detect and monitor treatment of Karposi's Sarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HPV 16-18 E6/E7 gene segment screening method-related PCR primer P3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR; primer; ss; gene screening method; cervical cancer; P3;
human papilloma virus 16.18; E6/E7 gene segment; HPV 16.18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15; DB 1; Length 20;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
Human; Kaposi's sarcoma; tumour; angiogenesis; tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 8 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Scor.
100.0%; Pred. No. 1...
0; Mismatches
                                                                                                                                                                                                                                                                                            В.
                                                                                                                                                                                                                                                                                            (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 9; Page 13; 38pp; English.
                                                                                                                                                                                                                                                                                                                                     Cornelissen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3704 CATGGTGGCCAGAGG 3718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL42402 standard; DNA; 20 BP
                                                                                                                                                                                                       23-JAN-2001; 2001EP-00200228.
28-SEP-2001; 2001EP-00203703.
28-SEP-2001; 2001US-0325722P.
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                                                                                                                                                              23-JAN-2002; 2002EP-00075264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-668396/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention
                                                                                                                                                                                                                                                                                                                                     Van Der Kuyl AC,
                                         Homo sapiens.
                                                                                  EP1225233-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JUN-2002
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                                                                                                                       24-JUL-2002
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Matches

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nucleotides. All 2' MOE

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Gaps

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Determining whether a treatment is effective in changing a status of a certain set of target cells in an individual comprises determining whether the sample comprises an expression product of at least one marker

Cornelissen M;

WPI; 2003-589342/56. Van Der Kuyl AC,

28-SEP-2001; 2001EP-00203703

(PRIM-) PRIMAGEN HOLDING BV.

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Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BBP alpha) gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions. The human and/or mouse C/BBP alpha RNA, and were analysed for their effect on C/BBP alpha mRNA levels by quantitative real-time PCR. The C/BBP family of proteins are a family of transcription factors which regulate the expression of a wide range of genes that control normal tissue development, cellular function, cellular proliferation and functional differentiation. C/BBP beta (also known as C/BPBI, LAP, TCFS, CRP2, NFL6, ILGDBP, NF-M, AGP/BBP and App/CBBP) primarily regulates hormone responsiveness and oxidative stress responses and is a mediator of IL-6 (interleukin-6) signalling. C/BBP beta is thought to be involved in carbohydrate metabolism, immunity, the Thi response, female fertility and gluconeogenic pathways. C/BBP beta is expressed in the liver, lung, spleen, kidney, brain, and testis, with the liver, it is also expressed at a higher control in the lung. It is also expressed at a higher
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        level in malignant ovarian tissue compared with normal ovarian tissue, and its expression in pancreas is upregulated in response to chronically elevated levels of glucose, indicating that it is involved in the impairment of insulin secretion in type II diabetes. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with C/EBP beta expression, such as cancer (particularly ovarian cancer), tumour formation, diabetes (particularly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         marker gene; tumour; Kaposi's Sarcoma; peripheral blood mononuclear cell; PBMC; expressed keratin 14; TIE 1; Salioadhesin; Siglec 1; angiogenesis; drug target; tag; SAGE library; KS3; KS4; RT-PCR; primer; ss.
                                                                                                                                                                                                                     Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/BBP) beta, useful in vitro for inhibiting expression of human or mouse C/EBP beta in cells/tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kaposi's sarcoma tag confirmation primer, SEQ ID No 46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 8 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  type II diabetes), infection, or inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 100.0%; Pred. No. 1.6 hes 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                             Example 17; Col 51-52; 69pp; English.
                                                                                                                                  Wyatt J;
14-JUN-2000; 2000US-00593711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC13379 standard; DNA; 20 BP.
                                           14-JUN-2000; 2000US-00593711.
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                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                  Butler MM,
                                                                                                                                                                              WPI; 2002-214451/27.
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                                                                                                                                Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC13379;
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treatment is effective in changing a status of a certain set of target cells in an individual. The method comprises obtaining a sample from an individual after initiation of the treatment, and determining whether the sample comprises an expression product of at least one marker gene. The marker gene and a proteinaceous molecule (which can bind to the protein derived from the marker gene of the invention) are useful for determining whether a treatment is effective in counteracting a tumour in an individual, especially Kaposi's Sarcoma. Peripheral blood mononuclear cell (PBMC) expressed keratin 14, TIE 1, Salioadhesin, or Siglec 1 sequences or a fully defined sequence given in the specification, or their analogues are useful as indicators for anglogenesis and for their analogues are useful as indicators for anglogenesis and for their analogues. The expression product of a gene comprising a marker gene of the invention is useful as a drug target. The compound is useful for preparing a medicament. This polymucleotide sequence represents a confirmation primer of a tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, antisense; lung dysfunction, nasal airway dysfunction, antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                           invention relates to a novel method for determining whether a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 15; DB 1; Length 20; 00.0%; Pred. No. 1.6e+03; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 3 C; 7 G; 2 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence of Kaposi's Sarcoma of the invention.
                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 46; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%,
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3704 CATGGTGGCCAGAGG 3718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ90612 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 CATGGTGGCCAGAGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-OCT-2003
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an national antiple associated with lung and/or has antiinflammatory, antiallargic, antiasthmatic, hypotenaive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an artificial matory steroid in a subject, for reducing levels of continuous or receptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of undiginant in subject; stissue, or treating bronchodilation, increasing levels of undiginal producing allarias or a consistent in a subject; stissue, or treating bronchodilation, increasing levels of undiginal producing increasing levels or consistion, lung inflammation in a subject or a consistion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                 treating ailments associated with impaired
                                                                                                                                                                                              respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                          ä
                          Aguilar
                       Sandrasagra A, Katz E, Pabalan J,
I, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 5854; 872pp; English.
                                                                                                                                                                 Pharmaceutical composition for
                                                                                                            WPI; 2003-229219/22.
                          Li Y, Sar
Tang L,
                                                                                                                                                                                                                                                      ubiquinone.
                          Nyce JW, 1
Miller S,
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Gaps ö 0.4%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 1.6e+03; ive 0; Mismatches 0; Indels Local Similarity 100. Les 15; Conservative Query Match Best Loc Matches

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ABD26842 standard; DNA; 20 BP (first entry) 29-JUL-2004 ABD26842;

AA278764-derived oligonucleotide SEQ ID 5854.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens

31-OCT-2002

WO200285309-A2

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

Katz E, Pabalan J, Aguilar D; Sandrasagra A, Ka L, Shahabuddin S; Tang L, Li Y, Miller S, Nyce JW,

WPI; 2003-093058/08

t 0 Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 5854; 763pp; English

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating control co e.g., lung, brain, heart, kidney, etc, prevent any unwanted effects due to it

Sequence 20 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 0 Other;

Gaps ò Query Match 0.4%; Score 15; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 1.6e+03; Matches 15; Conservative 0; Mismatches 0; Indels

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2165 CCCCACCCAGCAGTG 2179

ठ g

18 ccccaccascasts 4

ADL59284 standard; DNA; 20 BP. RESULT 1503 ADL59284

03-JUN-2004 (first entry)

ADL59284;

Human ESM-1 antisense oligonucleotide seqid 1533.

cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1stated disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se. nucleotides"

nucleotides'

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The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule.1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                          /mod_base= OTHER
/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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00.0%; Pred. No. 1.6e+03;
ve 0; Mismatches 0; Indels
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/note= "OTHER= 2'-methoxyethyl (2'-MOE)
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/note= "OTHER= 2'-methoxyethyl (2'-MOE)
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                                                                                           Location/Qualifiers
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/*tag= c
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/*tag=
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/*tag=
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Best Local Similarity
Matches 15; Conserv
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                                                                       /mod_base= OTHER
/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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                             Location/Qualifiers
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/mod_base= OTHER
/note= "OTHER= 2'
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Gaps

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gene therapy; endothelial specific molecule-1; ESM-1; SRM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.

cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective;

Human ESM-1 antisense oligonucleotide segid 1553.

(first entry)

03-JUN-2004

ADL59304;

/mod_base= OTHER /note= "OTHER= phosphorothioate backbone. All cytidine residues are 5-methylcytidines"

Location/Qualifiers

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modified base

sapiens

Homo

nodified base

/*tag= a /mod_base= OTHER /note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"

/mod_base= OTHER /note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"

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/*tag=

16. .20

modified base

19-AUG-2003; 2003WO-US025833 19-AUG-2002; 2002US-0404495P

WO2004021978-A2

18-MAR-2004

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The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule. (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
                                cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1selated disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
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/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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100.0%; Pred. No. 1.6e+03;
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 Human ESM-1 antisense oligonucleotide segid 1710.
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/mod_base= OTHER
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100.0%; Pred. No. 1.6e+03;
1ve 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 1.6 Matches 15; Conservative 0; Mismatches

3035 TAAAGCTATTTATGG 3049

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ADL59304 standard; DNA; 20 BP

RESULT 1506 ADL59304 ID ADL5 XX

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New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular

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Griggs

Weinstein EJ,

WPI; 2004-248358/23

(PHAA) PHARMACIA CORP

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The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, antisense oligonucleotide that can be used to modulate expression of endothelial specific molecule-1 (ESM-1).
                                                                                                                                                                       gene therapy; endothelial specific molecule-1; ESM-1;
ESM-1 related disorder; diabbtes; cancer; ischaemla; reperfusion injury;
angiogenic disorder; immunological disorder; cardiovascular disorder;
neurological disorder; antisense technology; ss.
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                                                                                                                                                         neuroprotective;
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/mod_bage= OTHER
/mode="OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
/fore= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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/mod_base= OTHER
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                             ADL59367 standard; DNA;
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RESULT 1507
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gene therapy; endothelial specific molecule-1; ESM-1;
ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury;
angiogenic disorder; immunological disorder; cardiovascular disorder;
neurological disorder; antisense technology; ss.
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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/notē= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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/note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                        Human ESM-1 antisense oligonucleotide segid 1518.
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                                                       ADL59269 standard; DNA; 20
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                                  RESULT 1508
                                             ADL59269
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The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disporder. This sequence represents an antisense oligonuclectide that can be used to modulate expression of endothelial specific molecule-1 (ESM-1).

Sequence 20 BP; 7 A; 1 C; 5 G; 7 T; 0 U; 0 Other;

Query Match

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Gaps

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0.4%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 1.6e+03; ive 0; Mismatches 0; Indels

DB 1; Length 20; 0.4%; Score 15;

Best Local Similarity 100. Matches 15; Conservative

Query Match

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Homo sapiens.
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                                                                                                                                                                                        chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammation; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
             Gaps
                                                                                                                                                                                                                                                                                                                                                                          /note= "phosphorothioate linkages and all cytidine
                                                                                                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1491
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            Indels
   Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                    residues are 5-methylcytidines
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16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "2'-0-methoxyethyls"
            Mismatches
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                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
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0
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                                 3035 TAAAGCTATTTATGG 3049
 100.08;
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                                             3 TAAAGCTATTTATGG 17
                                                                                                          ADM15304 standard; DNA; 20
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/mod_base=
                                                                                                                                                 (first entry)
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            15; Conservative
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                               modified base
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                                                                                                                                                                                                                                                                                                    sapiens
                                                                                                                                                 01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-APR-2004
                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gierse JK;
                                                                                                                            ADM15304;
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antidiabetic, immunomodulator, cardiant, neuroprotective, antiantlammatory, neuroprotective, nootropic, antiantitic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGBS-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diagnosing or predicting susceptibility to Crohn's disease in individual, comprises determining presence or absence of disease-predisposing haplotype comprising JWI variant allele and/or 268S allele at NOD2/CARDIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Crohn's disease; NOD2/CARD15 locus; single nucleotide polymorphism; SNP; autoimmune disease; psoriasis; ulcerative colitis; myasthenia gravis; autoimmune gastritis; Type I diabetes; 88; primer.
inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 15; DB 1; Length 20;
100.0%; Pred. No. 1.6e+03;
ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Targan SR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 12 A; 7 C; 1 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NOD2/CARD15 sequencing primer #9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.4
Best Local Similarity 100.
Matches 15; Conservative
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TARGAN S R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TAYLOR K D. ROTTER J I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2004076960-A1
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Sequence 25 BP; 0 A; 2 C; 0 G; 23 T; 0 U; 0 Other;

732

ô The useful for diagnosing or predicting susceptibility to a variety of autoimmune diseases such as Crohn's disease, psoriasis, ulcerative colities, mysathenia gravis, autoimmune gastritis and Type I diabetes. present sequence represents a primer used to sequence the nucleotide sequence of NOD2/CARD15. Gaps gene typing; polymorphic microsatellite loci; PML; disease; disease predsisposition; microsatellite marker; prion disease; eystic fibrosis; mailgnant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT-blue-vector; cow; ö 0.4%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 1.6e+03; ive 0; Mismatches 0; Indels Cow prion protein microsatellite locus primer #68. Seguence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other; microsatellite; PCR; primer; ss 1874 TGGAGGAGCTCTTCA 1888 ВР 2 TGGAGGAGCTCTTCA 16 ADO81056 standard; DNA; 25 (first entry) 15; Conservative Local Similarity DE10236711-A1. 29-JUL-2004 Bos taurus, ADO81056; Query Match 1511 Matches Best 8 88888888 ò

Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons. 09-AUG-2002; 2002DE-01036711. 09-AUG-2002; 2002DE-01036711 Preuss S, (UYHO-) UNIV HOHENHEIM WPI; 2004-215730/21. Geldermann H, 26-FEB-2004.

Example 3; Page 28; 64pp; German.

The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using a template a DNA sample containing at least one segment of (I); and cetermining the length of the resulting amplicon(s). Also described are: a method of determining (M3) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a disease and for prediagnosis of such diseases, especially prion diseases and for prediagnosis of such diseases, especially prion diseases but also expert (fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, continually less expensive than known methods based on sequencing. This equence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus.

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Gaps

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Score 15; DB 1; Length 25; Pred. No. 2e+03; 0; Mismatches 5; Indels

0.4%;

Query Match 0.4 Best Local Similarity 78.3 Matches 18; Conservative

Sequence 25 BP; 0 A; 2 C; 0 G; 23 T; 0 U; 0 Other;

The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and cetermining the length of the resulting amplicon(8). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and predisposition to a disease, associated with a gene that include PML. The method is used to identify microsatellite markers, in a disease and for prediagnosis of such diseases, especially prion diseases and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormones or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus.

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                       Gaps
                                                                                                                                                                                               gene typing; polymorphic microsatellite loci; PML; disease; adisease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT'-blue-vector; cow;
                       ;
0
0.4%; Score 15; DB 1; Length 25; 78.3%; Pred. No. 2e+03;
                      Indels
                      5;
                                                                                                                                                                            Cow prion protein microsatellite locus primer #73.
                      0; Mismatches
                                           3262 TATTTTATTTGCTTTGTCCTTTT 3284
                                                        3 TTTTTTTTTTTTTTTCCTTTT 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 28; 64pp; German.
                                                                                                                                                                                                                                                                                                                                                                                   Han Y;
                                                                                                                                                                                                                                         microsatellite; PCR; primer; ss
                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                         09-AUG-2002; 2002DE-01036711,
                                                                                                                                                                                                                                                                                                                                            09-AUG-2002; 2002DE-01036711.
                                                                                                                 ADO81061 standard; DNA; 25
                                                                                                                                                         (first entry)
                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                     Preuss S,
                                                                                                                                                                                                                                                                                                                                                                 (UYHO-) UNIV HOHENHEIM.
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-215730/21
           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                  Geldermann H,
                                                                                                                                                                                                                                                                                DE10236711-A1
                     18;
                                                                                                                                                         29-JUL-2004
                                                                                                                                                                                                                                                                                                     26-FEB-2004.
                                                                                                                                                                                                                                                              Bos taurus.
                                                                                                                                     ADO81061;
   Query Match
                                                                                           RESULT 1512
                      Matches
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vivlemore401-10.rng

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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers pecifically priming and allowing amplification of each Vbeta gene, vbetaRNA or cDNA. The kit is useful for diagnosing organ transplant respection and diagnosing and treating T-cell associated diseases including autoimmune disease, degenerative nervous system diseases, infectious diseases including autoimmune disease, degenerative nervous system diseases, and neoplastic disease, Autoimmune diseases include Addison's disease. Carff versus host disease. Autoimmune diseases include Addison's disease, atrophic gastritis. Degenerative nervous system diseases include multiple atrophic gastritis. Degenerative nervous system diseases include multiple in hypersensitivities such as those present in allergies, Type II hypersensitivities such as those present in callergies. Type II hypersensitivities such as those oused by condpasture's syndrome and Type IV hypersensitivities such as those caused by the yeast genus Candida, parasitic infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by Mycobacterium. Neoplastic diseases include lymphopyoliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                   human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; HIV; kungal infection. Candida; parasitic infection; schistoseme; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kit for diagnozing and treating T-cell associated diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 971; 164pp; English.
                                                                                                                                                                                                                                                                                  Human Vbeta gene repeat sequence #567.
3262 TATTTTATTTGCTTTGTCCTTTT 3284
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                                                                                                                                               ADH70777 standard; DNA; 18
                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      breast cancer; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2002150891-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                  receptor; flt-1; flk-1;
                                                                                                                                                                                                                                                                                                                                Vascular endothelial growth factor receptor; VEGF receptor; flt-1; fl)
KDR; hammerhaed ribozyme; hairpin ribozyme; cleavage;
tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                              Gaps.
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                             Score 14.8; DB 1; Length 18;
Pred. No. 1.5e+03;
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Pred. No. 1.5e+03;
                                                            Indels
                                                                                                                                                                                                                                                                                                    Mouse flk-1 VEGF receptor hairpin ribozyme substrate #41.
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G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 18 BP; 1 A; 6 C; 7 G; 0 T; 4 U; 0 Other;
                                                              0; Mismatches
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                                                                                            2824 ATATATACATATATAT 2841
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Sequence 18 BP; 8 A; 0 C; 1
                               0.4%;
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                                              88.9%;
                                                                                                                         1 ATATATATATATATAT
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                                                                                                                                                                                                                                                                     28-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                    foetal liver kinase 1; ss.
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                                              Local Similarity 88.9
les 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9715662-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-OCT-1995;
                                                                                                                                                                                                                                      AAX73494;
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                               Query Match
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Matches
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28-JUL-1999

AAX70271;

1515

RESULT

25-OCT-1996; 26-OCT-1995; 11-JAN-1996;

01-MAY-1997.

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AAV02501-V02522, AAV02524-V02584, AAV02586-V02592 and AAV02594-V02616 are DNA fragments used in an assay to determine novel transcriptional activators. The method involves the production of transcriptional activators comprising of a DNA-binding group and a 6-25 amino acid peptide that is covalently bonded to the DNA binding group and does not cepresent a fragment of a natural transcription activator. Protein-C protein interactions are identified in the assay by fusing a DNA-binding domain to a library of DNA fragments and introducing this and a fusion of carget protein and a polypeptide containing a region of Gal4 which interacts with Gal11P into a cell containing dal11P and identifying members of the library that interact with the target from activation of caranscription. Such constructs are used to activate transcription in a cell, e.g. recognizing a site close to a selected therapeutic gene).

CC Transcription can be activated without blocking other transcription and activators. They probably act by interacting with a component of the RNA polymerase II holoenzyme, Gal11, the strongest known yeast activator. Death and activators. They provides a more sensitive assay allowing detection of even weak contains a contains and a provides a more sensitive assay allowing detection of even weak contains and a processed to a more sensitive assay allowing detection of even weak contains and a contains a more sensitive assay allowing detection of even weak contains and a conta
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sequence; Gal4; transcriptional activator; RNA polymerase;
otein interaction; gene therapy; therapeutic; holoenzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New transcriptional activator containing DNA binding domain bound to peptide - useful for controlling gene expression, especially in gene therapy, and in protein-protein interaction assays, does not inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein-protein interactions. Such activators do not create toxicity problems even when overexpressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transforming growth factor beta-1 antisense oligonucleotide N37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 26; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2698 CTTCCCACCCTGCCCTC 2715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other transcription activators.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18
                       Protein-protein interaction; Gall1; DNA binding domain; 88
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97US-00017016.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wu Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            (HARD ) HARVARD COLLEGE
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                                                                                                                                                                      WO9744447-A2
                                                                                                                                                                                                                                                                                                                                         03-MAY-1996;
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                                                                                                                                                                                                                           27-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ptashne M,
                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endochelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or focetal liver kinase 1 (fit-1) (eg. tumour angiogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression
                                                                                                                                                                                                                                                                                                           Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention describes nucleic acid molecules which modulate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                   Human flt1 VEGF receptor hairpin ribozyme substrate #39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Escopedo J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 5 C; 5 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stinchcomb D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 93; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1391 TCAACCTGCTGGCCCCT 1408
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UVAACCUGCUGGGAGCCU 18
                                                                          AAX70271 standard; RNA; 18 BP.
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96US-00584040.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV02562 standard; DNA; 18
                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pavco P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-259017/23
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Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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Gaps

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04-AUG-1998

AAV02562; RESULT 1516
AAV02562
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AC AAV02562
AC AAV02562
DT 04-AUG-1
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Synthetic

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Treating rheumatoid arthritis of an individual comprising expressing an immunologically recognizable portion of a cytokine to induce formation of anti-CK immunoglobulins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bovine; leukocyte interferon; IFN; bovine IFN-beta; porcine IFN-alpha1; porcine IFN-beta1; porcine IFN-gamma; feline IFN-beta1; rabbit IFN-gamma; non-human mammalian interferon; viral infection; malignant condition; immunosuppresed condition; immunodeficient condition; respiratory complex; interferon-alpha1; IFN-alpha1; ss.
                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method of treating rheumatoid arthritis, involving expressing in the individual an immunologically recognisable portion of a cytokine such that anti-cytokine immunoglobulins are produced, neutralising the activity of an endogenous cytokine and therefore treating rheumatoid arthritis. The present sequence is a PCR primer for the rat beta-actin cDNA described in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Porcine interferon-alphal (IFN-alphal) expression associated DNA #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 4 A; 8 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                         (TECR ) TECHNION RES & DEV FOUND LID
                                                                                                                                                                                                                                                                                                                                     Example; Page 21; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    550 CTGCCGGCCAACCAGACG 567
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82US-00438128.
87US-00104461.
91US-00749371.
                       05-FEB-2001; 2001WO-IL000117
                                                                 07-FEB-2000; 2000US-00498625
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Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Goeddel DV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-689760/74.
                                                                                                                                                                                                  WPI; 2001-476277/51.
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01-NOV-1982;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-AUG-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Capon DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS54009;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                        Karin N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1519
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV48412-84 represent antisense oligonucleotides directed against transforming growth factor beta-1 (TGP beta-1). The oligonucleotides exemplify the invention. The specification describes oligonucleotides (that contain B-30 nucleotides, which contain at most 8 nucleotides that can each form three hydrogen bonds to cytosine; do not contain four consecutive nucleotides able to form three H-bonds each to four consecutive cytosines; do not contain two sequences of three consecutive nucleotides each able to form three H-bonds to three consecutive cytosines, and the ratio between residues able to form two H-bonds each colloquelectides are used to modulate expression of genes, particularly the groundleotides are used to modulate expression of genes, particularly the groundleotides are used to modulate expression of genes, particularly the groliferation of primary cell cultures (e.g. bone marrow stem, liver or kidney cells, osteoclasts, osteoblasts and/or keratinocytes). The coligonucleotides can also be used to analyse function of proteins (by altering their expression or activity) and therapeutically, e.g. in cases of cancer or (targeting TGF) for stimulating the immune system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                    Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rat; beta-actin; rheumatoid arthritis; DNA vaccine; cytokine; self-tolerance; inflammation; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 18; 38.9%; Pred. No. 1.5e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 0 A; 3 C; 15 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                   (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Fig 3a; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat beta-actin cDNA PCR primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2920 GGCCGGCCGTGGGGGG 2937
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                                                                                                                                                                                                                                                                                                                Schlingensiepen K, Brysch W;
                                                                                                                                                                                                                            97EP-00101531
                                                                                                                                                                               97EP-00101531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-400910/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200157056-A1
                                              Homo sapiens
                                                                                                                                                                               31-JAN-1997;
                                                                                                                                                                                                                         31-JAN-1997;
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                                                                                                                                   05-AUG-1998
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AAH73453;

RESULT 1518

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Query Match

Best Loc Matches

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Gaps

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The invention describes a purified and isolated nucleic acid (I) encoding a non-human mammalian interferon (IFN) chosen from bovine leukocyte IFNs, bovine IFN-betal 1, 2 or 3, porcine IFN-alphal, IFN-betal, IFN-gamma, feline IFN-betal or rabbit IFN-gamma. A microorganism or cell culture transformed with an expression vector comprising nucleic acid encoding the non-human mammalian interferon is useful for producing a polypeptide consisting of the amino acid sequence of a non-human mammalian IFN. The non-human animal IFNs are useful in the prophylactic or therapeutic treatment of non-human animals, in particular for viral infections, and malignant, and immunosuppressed or immunodeficient conditions. Bovine IFNs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represents an oligonuclectide used to incorporate a peptide encoding an ATG translation initiation codon, DNA encoding which is incorporated into a plasmid for expressing porcine interferon-alpha! (IFN-alpha!)
including bovine, porcine, feline, and rabbit interferons, useful for treating viral, malignant, and immunosuppressed or immunodeficient
                                                                                                                                                                                                                                                                                                                                                                                                                                                         find use in treating respiratory complex in cattle. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 4 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                English.
                                                                                                             Disclosure; Col 21; 42pp;
                               treating vi
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0.4%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 1.5e+03; ve 0; Mismatches 2; Indels 512 18 GTACACGCTGGACGTGCT 88.9%; Gracacacragacacacr Local Similarity 88.5 hes 16; Conservative 495 Query Match g 8

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Gaps

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Drosophila genghis khan, gek, PCR primer GK3 ABK13419 standard; DNA; 18 (first entry) Drosophila melanogaster 23-APR-2002 ABK13419; RESULT 1520

BP

rot; insecticide; MYST; acaricide; Fruit fly; ss; rotkehlchen; rot; in genghis khan; gek; GK3; PCR; primer

WO200200864-A2.

03-JAN-2002

08-JUN-2001; 2001WO-EP006505.

27-JUN-2000; 2000EP-00113527.

(AVET) AVENTIS CROPSCIENCE GMBH.

Benting J, Gunkel N;

Luemmen P,

WPI; 2002-130888/17.

Pankratz MJ, Zinke I,

Novel isolated DNA molecule encoding protein having biological activity of histone acetyltransferase which is useful for screening histone acetyltransferase inhibitors that serve as insecticides and acaricides.

Example B; Page 21; 61pp; English.

The invention relates to an isolated DNA molecule comprising a DNA sequence which encodes an insect histone acetyltransferase (HAT a member of the MYST family which regulates food uptake) and is either the Rotkehlchen (rot) gene or the rot CDNA, or their fragments, derivatives or allelic variants. Also included are a vector comprising the nucleic acids, a eukaryotic cell harbouring the vector or nucleic acids and an

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biochemical activity of HAT when compared with the non-treated control biochemical activity of HAT when compared with the non-treated control brotein in presence of suitable substrate, buffer and assay conditions. The vector is useful for the recombinant production of ROT. ROT is useful for the biochemical or structural characterisation of the potential inhibitors of the encoded protein. The inhibitor, in appropriate chemical compositions, is useful for an insect controlling method based on specific inhibition or sufficient reduction of activity of the native target protein (Rotkehlchen (ROT) protein which is a HAT that belongs to the so called MAST family of HAT) in an insect. ROT protein is useful as insecticide or acaricide. The inhibitor has agrochemistry, veterinary and pharmaceutical applications. The present sequence is a PCR primer used to isolate sequence encoding the ROT protein, in this case being derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotides, useful for the modulation of Smad6 expression in the treatment or prophylaxis of e.g. cardiovascular disease, are targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, Smad6 protein, antisense, cardiovascular disease, infection, inflammation, cancer, therapy, phosphorothioate backbone, ss.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1; Length 18;
Pred. No. 1.5e+03;
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/note= "OTHER = Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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/note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Smad6 antisense oligonucleotide, ISIS #28571.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 4 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 from the adjacent gene genghis khan, gek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2068 GCGCCTTTCGAGCAGTAC 2085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GCGCTTATCGAGCAGTAC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD36203 standard; DNA; 18
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD36203;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1521
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                   88999999999999988
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16; Conservative
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                                                                                           RESULT 1523
 Matches
                                                                                                          ADB80921
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to novel remedies for the treatment of diseases containing as an active component an inhibitor of the expression or activity of hypothalamus-expressed G-protein coupled receptor 901 and for treatment of cibophobia containing as an active component a potentiator of the expression or activity of G-protein coupled receptor 901. The diseases which can be treated include obsesty, diabetes and hyperlipaemia, and cibophobia (anorexia nervosa). The present oligonucleotide was used in an example from the invention
                                                 The invention relates to an antisense oligonucleotide targetted to a nucleic acid molecule encoding human Smad6 protein, which specifically hybridises with the nucleic acid and inhibits its expression. Antisense compounds of the invention are used for inhibiting the expression of Smad6 in cells and tissues in the treatment of a disease or condition associated with Smad6 such as cardiovascular disease, cancer, infection and inflammation. They are also useful in the disgnostics, as research reagents, in kits and in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human Smad6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inhibitors of expression or activity of G-protein coupled receptor 901 for treatment of lifestyle-related diseases and cibophobia.
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; anorectic; antidiabetic; antilipemic; hypothalamus;
G-protein coupled receptor 901; obesity; diabetes; hyperlipaemia;
cibophobia; anorexia nervosa; 88.
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                                                                                                                                                                                                            Query Match 0.4%; Score 14.8; DB 1; Length 18; Best Local Similarity 88.9%; Pred. No. 1.5e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 A; 0 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                      Sequence 18 BP; 5 A; 1 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Taiji M;
nucleic acid molecule encoding Smad6.
                           16; Page 90; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                           6xHis-tag linker oligonucleotide #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yamanaka M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 68; 91pp; Japanese.
                                                                                                                                                                                                                                                                    185
                                                                                                                                                                                                                                                                                            1 GGGAGTTGACGAAGATGG 18
                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SUMU ) SUMITOMO PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-OCT-2002; 2002WO-JP010250
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                                                                                                                                                                                                                                                                    168 GGGAGATGACGAAGACGG
                                                                                                                                                                                                                                                                                                                                                              ACC70855 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-354886/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sugaru E,
                                                                                                                                                                                                                                                                                                                                                                                        ACC70855;
                           Example
                                                                                                                                                                                                                                                                                                                                      RESULT 1522
                                                                                                                                                                                                                                                                                                                                                    ACC70855
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Score 14.8; DB 1; Length 18; Pred. No. 1.5e+03;

0.4%;

Query Match Best Local Similarity

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The invention relates to novel remedies for the treatment for anorexia and lifestyle related disorders, comprising a substance that prevents expression or function of a polypeptide having a 300 or 345 residue amino acid sequence, given in the specification. The invention further relates to a nucleic acid comprising a 1038 or 1324 nucleotide sequence, given in the specification. The novel remedies have anabolic activity and can be used to treat disorders by gene therapy or cell therapy. The remedies can be used in the treatment of anorexia and lifestyle related disorders such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                              primer, lifestyle related disorder, anabolic activity, gene therapy; cell therapy; eating disorder; obesity; diabetes; hypertension; ss; anorexia; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hybridisation oligonucleotide 74 used to analyse genomic DNA region.
                                                                                                                                                                                                                                                                                                                                                                                         Anorexia / life-style related diseases related primer, SEQ ID No 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treatment for anorexia, eating disorders, obesity, diabetes and hypertension by preventing expression or function of a polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Taiji M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 7; Page 87; 91pp; Japanese.
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1351 ATGGAGATGATGAGATG 1368
                                                       1 Argergargargarg 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SUMU ) SUMITOMO PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-DEC-2001; 2001JP-00397523.
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                                                                                                                                                                                                         ADB80921 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB54536 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                              04-DEC-2003 (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-618060/58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003055507-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-2003
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ADB54536/C
XX
AC ADB54536
XX
XX
D4-DEC-2
XX
D5
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D7
D6
Hybridie
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Ξ, Carr

probe.

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The invention relates to a novel modified bacterial enzyme carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations compared with the non-modified parochial enzyme. The alterations cause a reduction or an elimination of one or more of T-cell epitope sequences, which act in the parental enzyme as MHC class II binding ligands and stimulate T-cells. The modified CPG2 enzyme and the CPG2 proteins have immunostimulant activity and may be used in enzyme therapy. The modified CPG2 enzyme may be used to induce an immune response in a human host, or as a therapeutic entity such as the gene classification of modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo. This polymucleotide sequence represents a mutagenic oligonucleotide used in the production of a modified CPG2 gene of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New modified bacterial enzyme carboxypeptidase G2 (CPG2) having substantially non-immunogenic or less immunogenic than any non-modified CPG2, useful for inducing an immune response in a human host.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.4.17.11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Carboxypeptidase G2 (CPG2) enzyme mutagenic oligonucleotide OL589.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 7 A; 3 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                   Hellendoorn K, Baker M, Williams S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 36; 52pp; English
                                                                                                                                                                                                                                        25-JAN-2002; 2002EP-00001778.
13-SEP-2002; 2002EP-00020634.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD44224 standard; DNA; 18 BP
                                                                                                                                                         27-NOV-2002; 2002WO-EP013351,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2754 TACCTTTTATGCAAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 TACCTTTTATGTAAACGG
                                                                                                                                                                                                                                                                                                                  (MERE ) MERCK PATENT GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16; Conservative
     Pseudomonas sp. RS-16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
Pseudomonas sp. RS-16.
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-513617/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                   WO2003045426-A1
                                                                                                                                                                                                             29-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mutagenic; ss
                                                                                                       05-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD44224;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1526
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD44224
ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel method for detecting and differentiating between colon cell proliferative disorders associated with at least one gene or its regulatory regions. The method comprises contexting a target nucleic acid in a biological sample obtained from the subject with at least one reagent or a series of reagents, where the reagent or series of reagents, distinguishes between methylated and non methylated CpG dinucleotides within the target nucleic acid. The molecules of the invention demonstrate cytostatic activity whilst the method may useful to describe and differentiating between colon cell proliferative disorders, including cancers such as colon adenoma and colon carcinoma. The PNA (peptide nucleic acid) oligomers are useful as probes for determining cytosine methylation state or single nucleotide polymorphisms. The current sequence is that of the hybridisation control colon carcinoma colon methylams. The current sequence is that of the hybridisation colon carcinoma colon methylams. The current sequence is that of the hybridisation colon carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting and differentiating between colon ceil proliferative disorders associated with a gene or its regulatory regions comprises contacting a target nucleic acid in a biological sample obtained from the subject with
                   colon cell proliferative disorder; non methylated CpG dinucleotide; cytostatic; cancer; adenoma; carcinoma; cytosine methylation state; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MTC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.4.17.11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                             Lesche R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Carboxypeptidase G2 (CPG2) enzyme mutagenic oligonucleotide OL573.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 1.5e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                Becker E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 2 A; 0 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Maier S, Nimmrich I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 36; Page 30; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1379 ACAAAAACATCATCAACC 1396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD44208 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                 27-FEB-2003; 2003WO-EP002035.
                                                                                                                                                                                                                                                                                                                                     27-FEB-2002; 2002EP-00004551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACAAAAACATCATCCCCC
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                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                             Burger M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schmitt A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-731620/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                            WO2003072821-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mutagenic; ss.
                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                          Adorjan P,
Rujan T, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          a reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Gaps

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2; Indels

0; Mismatches

2771

Synthetic

RESULT 1525

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Matches

Score 14.8; DB 1; Length 18; Pred. No. 1.5e+03;

0.4%;

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This invention describes a novel system of screening for ligands of the bilayer membrane coupled receptor (GPCR). The method comprises a lipid bilayer membrane in which a fusion protein comprising target GPCR and Galpha i. Or Galpha is or Galpha is or Galpha 2 is embedded. The invention also discloses a second screening system where an orphan GPCR (G protein comjugation seven-transmembrane-type receptor) is used to screen compound having agonist and/or antagonist activity for the GPCR and to screen low molecular non-peptide ligands. The screening is rapid and favourable.
                                                                                                                                                                                  Screening ligands for G protein coupled receptor comprises lipid bilayer membrane containing embedded fusion protein comprising target G protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GPCR; G protein-coupled receptor; C-PLACE 1003238; ss; G16 alpha; G12 alpha; uropathic; gynaecological; GDP/GTP exchange reaction; urinary tract disease; placental disease; tonsil disease; Hisx6; linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ligand screening system comprising a component which is a lipid bilayer membrane that contains C-PLACE1003238 and a region concerned in binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 1.5e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 5 A; 0 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 13; 28pp; Japanese.
                                                                                                                                                                                                                                                                              Example 1; SEQ ID NO 12; 29pp; Japanese.
                                                                                                                                                                                                                                       coupled receptor and G alpha protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1351 ATGGAGATGATGAAGATG 1368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Argercarcarcarcars 18
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18-JAN-2002; 2002JP-00010871.
                                            18-JAN-2002; 2002JP-00010871.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-FEB-2002; 2002JP-00034569
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG36975 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 88.9
Matches 16; Conservative
                                                                                         (SUMU ) SUMITOMO SEIYAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SUMU ) SUMITOMO SEIYAKU
                                                                                                                                      WPI; 2003-819838/77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-014845/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hisx6 linker #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP2003232790-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG36975;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel modified bacterial enzyme carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations compared with the non-modified parochial enzyme. The alterations compared with the non-modified parochial enzyme. The epitope sequences, which act in the parental enzyme as MHC class II binding ligands and stimulate T-cells. The modified CPG2 enzyme and the CPG2 proteins have immunostimulant activity and may be used in enzyme therapy. The modified CPG2 enzyme may be used to induce an immune response in a human host, or as a therapeutic entity such as the gene manufacture of a modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, immunogenicity than any non-modified parental enzyme when used in vivo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 is polynucleotide sequence represents a mutagenicity to CPG2 in vivo.
the production of a modified CPG2 gene of the invention.
                                                                                                                                                                                                                                                                                                                                                                          New modified bacterial enzyme carboxypeptidase G2 (CPG2) having substantially non-immunogenic or less immunogenic than any non-modified CPG2, useful for inducing an immune response in a human host.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and for vaccination of patients to reduce immunogenicity to CPG2
This polynucleotide sequence represents a mutagenic oligonucleoti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88; screening; human; G protein coupled receptor; GPCR; lipid bilayer membrane; fusion protein; G-alpha 16; G-alpha 12; G-alpha S2; orphan GPCR; G protein conjugation seven-transmembrane-type receptor; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 5 A; 3 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                   Williams S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 37; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2754 TACCTTTTATGCAAAAGG 2771
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                                                                                                                                      29-NOV-2001; 2001EP-00128519.
25-JAN-2002; 2002EP-00001778.
13-SEP-2002; 2002EP-00020634.
                                                                                           27-NOV-2002; 2002WO-EP013351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6XHIS tag associated primer.
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                                                                                                                                                                                                                                       (MERE ) MERCK PATENT GMBH
                                                                                                                                                                                                                                                                                 Hellendoorn K, Baker M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 88.9
les 16; Conservative
                                                                                                                                                                                                                                                                                                                                WPI; 2003-513617/48.
WO2003045426-A1
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                                            05-JUN-2003
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ADG37249
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Matches

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The invention relates to a screening system of a ligand with respect to C -PLACE1003238 (a GPCR), where a ligand-receptor interaction promotes estivity of GDP or GTP exchange reaction of G-protein subunits comprises, a component which is a lipid bilayer membrane that contains a polypeptide having a region which is concerned in binding with guanine nucleotide in

JP2003210192-A.

29-JUL-2003

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Gaps

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of nucleic acids of HBV and/or HIV and/or HCV. The present sequence is used in the exemplification of the invention.

Sequence 18 BP; 8 A; 2 C; 7 G; 0 T; 1 U; 0 Other;

SXSS

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Gaps

; 0

Indels

Length 18;

Score 14.8; DB 1; Pred. No. 1.5e+03;

0.4%; 88.9%;

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G protein-coupling receptor (GPCR) of the G-protein alpha (G16 alpha or G12 alpha) subunit that belongs to the G1 family. Also included are screening a ligand (involving comparing the effect of effector when interacted with ligand in presence or absence of the test material), producing a prophylactic and therapeutic agent of diseases of the urinary tract, placenta or tonsil (involving mixing the effector and a carrier), identifying a marker substance of the disease in the urinary tract, placenta or tonsil (involving comparing the presence of ligand identified in the biological sample derived tonsils obtained from patients and normal humans), diagnosing the peptide which consists of amino acids 12-36 of C-placenta or consil and an antibody recognising the peptide which consists of amino acids 12-36 of C-placental susful in treating and preventing the disease in the tissue of urinary organ, placenta or tonsil. Hisxé fusion proteins were constructed with human G16 alpha and G12 alpha for use in the construction system. The present sequence is a Hisxé linker used in the construction of the fusion proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Capture oligonuclectide composition useful for detection of hepatitis B virus (HBV), comprising polynuclectide having HBV-complementary sequence which is immobilized on solid support.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; capture oligonucleotide; HBV; HIV-1; HCV; donated blood screening.
                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1; Length 18;
Pred. No. 1.5e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 5 A; 0 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Getman DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HBV specific molecular beacon target #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Linnen JM, Kolk DP, Dockter JM,
Ho-Sing-Loy M, Stringfellow LA;
                                                                                                                                                                                                                                                                                                                                                                                                                        1351 ATGGAGATGATGAAGATG 1368
                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 Argercarcarcarcars 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADJ53735 standard; RNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                    Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003106714-A1.
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Microsatellite markers based on hypervariable genomic fragments, from
Triticum aestivum (wheat) or the tribe Triticeae, consist of a sequence
tagged site (STS), defined by 2 specific primers (of mean size 17-23
bases) that flank a microsatellite sequence at both ends, which can be
microsatellites are n-fold tandem repeats (n = 10 or more) of di-, tri-
cor tetra-nucleotide sequences, combination microsatellite sequences or an
imperfect sequence in which individual bases are mutated. The
cor tetra-nucleotide sequences, combination microsatellite sequences or an
imperfect sequence in which individual bases are mutated. The
cor tetraploid forms of wheat and for genetic mapping or labelling of
tetraploid forms of wheat and for genetic mapping or labelling of
corrected forms of wheat and for genetic mapping or labelling of
monogenic and polygenic properties, and for their selection; for
cor nalysing relationships and identification and plant growth. The markers can
differentiate between almost all European wheat lines and show a higher
cor degree of DNA polymorphism than known probes for the wheat genome. They
can be detected by PCR, so large numbers of samples can be analysed
can be detected by PCR, so large numbers of samples can be analysed
can be detected by PCR, so large numbers of samples can be analysed
colymorphisms are stably inherited so can also serve as genetic markers.
CAAT77003-22 and AAT77353-716 are primer pairs that define the
corrected properties and can be defined the markers.
                                                                                                                                                                                                                                                                                                                                                                                                                          Microsatellite marker, hypervariable genomic fragment, Triticum aestivum, wheat, Triticeae, sequence tagged site, STS, primer, PCR, amplify, polymorphism, genetic analysis, hexaploid, tetraploid, mapping, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primers for STS micro:satellite markers for wheat and related species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful for genetic mapping, analysis and labelling etc. of wheat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PFLA-) INST PFLANZENGENETIK & KULTURPFLANZENFOR.
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                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                             Wheat microsatellite WMS179 right primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ganal M;
                                                                                      925 TTCCTGTTCATCCTGGTG 942
                                                                                                                                18 rrccrcrrcarcard 1
                                                                                                                                                                                                                                               BP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95DE-01025284.
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                                                                                                                                                                                                                                            AAT77620 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                      11-SEP-1997 (first entry)
Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-053731/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                           AAT77620;
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0.4%; Score 14.8; DB 1; Length 19;

Query Match

The invention relates to a capture oligonucleotide composition comprising an hepatitis B virus (HBV)-complementary sequence polynucleotide immobilised to a solid support. The composition is useful for detecting nucleic acids of HBV and/of HTV-1 and/or HCV in biological sample such as blood, serum, plasma or other body fluid or tissue to be tested. The composition can be used either in diagnostic application or for screening donated blood and that products or other tissues that may contain infectious particles. The composition facilitates detection of very low levels of HBV nucleic acids. The composition allows selective detection

Claim 22; SEQ ID NO 129; 112pp; English.

PCR primer; chromosomal abnormality; abnormality detection; leukaemia; lymphoma; carcinoma; adenocarcinoma; sarcoma; glioma; neuroblastoma; medullablastoma; malignant melanoma; malignant neoplastic condition; ss.

Homo sapiens WO9824928-A2

Synthetic

11-JUN-1998

Primer ALL1ENL:4195L19 for abnormality detection.

25-SEP-1998 (first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genetic detection of primary adhalinopathies - using nucleic acid probes which bind to mutant adhalin genes but not the wild type gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Adhalin gene fragment where muscular dystrophy causing mutation occurs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; adhalin gene; dystrophin-associated protein; muscular dystrophy; detection; mutation; primary adhalinopathy; Duchenne-like autosomal recessive muscular dystrophy; probe; ds.
                                           Gaps
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88.9%; Pred. No. 1.6e+03;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 3 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           769,
                                                                                                         404 AGTGGAGCCTGGTCATGG 421
                                                                                                                                                     18 AGTGGATGCTGGTCATGG 1
                                                                                                                                                                                                                                                                                                                                                           BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                        16; Conservative
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Best Local Similarity
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Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                            AAV22618;
                                                                                                                                                                                                                                                                              RESULT 1531
AAV22618
XX
AAV22618
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DT 08-JUL-XX
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HUMMAN;
KW HUMAN;
KW HUMAN;
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KW DUCHENN
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Detection of chromosomal abnormalities - by subjecting patient sample nucleic acids to a multiplex molecular amplification procedure using primers specific for characteristic nucleic acid sequence.

96DK-00001401.

06-DEC-1996;

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Pallisgaard N, Hokland PALL/) PALLISGAARD N.

WPI; 1998-333344/29.

Claim 73; Page 105; 126pp; English

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This sequence represents a primer used in the method of the invention for the detection of the presence or absence of chromosomal abnormalities, cach abnormality being associated with a condition in a subject and each being defined by at least one characteristic nucleic acid sequence. The method comprises: (a) obtaining a sample of nucleic acids derived from a subject which may harbour one of the chromosomal abnormalities; (b) subject which may harbour one of the chromosomal abnormalities; (b) procedure, where a number of the characteristic sequences; (b) procedure, where a number of the presence and/or absence of a namplicon characteristic of the abnormal sequences to detect the presence of characteristic of the abnormal sequences to detect the presence of characteristic of the abnormal abnormalities; where the MMA concessing the presence of detecting the presence of detecting the presence of the primers defining an end of at least one characteristic nucleic acid sequence, and where at least one of the primers defines the first end of at least two characteristic nucleic acid sequences each being determined in their opposite ends by MDP selected from the remainder of the MDP The methods can be used for detecting chromosomal abnormalities
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   associated with diseases including numerous leukaemia's, lymphoma's, carcinoma's, adenocarcinoma's, sarcoma's, glioma's, neuroblastoma's, medullablastoma, malignant melanoma, and malignant neoplastic conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 0 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
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nes 16; Conservative
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18

TGCTCAACGTCACCTCTG

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AAV41059 standard; DNA; 19

RESULT 1532

AAV41059

AAV41059 ID AAV4 XX AC AAV4 XX

vivlemore401-10.rng

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The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polynuclectide and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allelespecific amplification assay for determining the identity of a nuclectide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate cancer. Early-stage diagnossis of prostate cancer relies on prostate specific antigen (PSA) dosage. However, the effectiveness of this is limited due to its of the organ. A need exits for both a reliable diagnostic procedure which would enable early-stage diagnosis, and for preventative and detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast; PCR primer; URA3; chromosomal rearrangement; double strand break;
                                                                                                                                                                                                                                                              Use of a prostate cancer associated gene and biallelic markers derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 14.8; DB 1; Length 19;
88.9%; Pred. No. 1.6e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                        Bougueleret L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 8 A; 2 C; 0 G; 9 T; 0 U; 0 Other;
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                                                                                                                                                                                        Chumakov I,
                                                                                                                                                                                                                                                                                                                      Claim 4; Page 369; 385pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3469 TATCTATATATATTT 3486
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                                                                                           97US-00996306.
98US-0099658P.
                                                        98WO-IB002133
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Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces cerevisiae.
                                                                                                                                                                                    Blumenfeld M,
                                                                                                                                                                                                                         WPI; 1999-405178/34.
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                                                                                                                                                   (GEST ) GENSET
                                                        22-DEC-1998;
                                                                                                              09-SEP-1998;
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                     01-JUL-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gabriel A;
                                                                                                                                                                                        Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF60558;
                                                                                                                                                                                                                                                                                   Erom it.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polynucleotide and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allelespecific amplification assay for determining the identity of a nucleotide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate cancer. Early-stage diagnosis of prostate cancer relies on prostate specific antigen (PSA) dosage. However, the effectiveness of this is limited due to its inability to discriminate between malignant and non-malignant affections of the organ. A need exits for both a reliable diagnostic procedure which would enable early-stage diagnosis, and for preventative and detection of prostate cancer, and the arisk of developing it in the future, and can also be used to determine therapies for the disease
                     PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate specific antigen; PSA; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen; PSA; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of a prostate cancer associated gene and biallelic markers derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                             Chumakov I, Bougueleret L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 19 BP; 2 A; 0 C; 7 G; 10 T; 0 U; 0 Other;
primer for PG1 biallelic marker 99-128-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer for PG1 biallelic marker 99-148-129.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 379; 385pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2315 GTCTGTGTGTGTGTGT 2332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Grargrargrargrar 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ01326 standard; DNA; 19 BP.
                                                                                                                                                                                                                                               98WO-IB002133.
                                                                                                                                                                                                                                                                                 97US-00996306
                                                                                                                                                                                                                                                                                                   98US-0099658P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 88.9
les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                         Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-405178/34.
                                                                                                                                                                                                                                                                                                                                      (GEST ) GENSET
                                                                                                                                Homo sapiens
                                                                                                                                                                   WO9932644-A2
                                                                                                                                                                                                                                             22-DEC-1998;
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                                                                                                                                                                                                                                                                                 22-DEC-1997;
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                                                                                                                                                                                                          01-JUL-1999.
                                                                                                              Synthetic.
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            Intron-based assay for detecting chromosomal rearrangements in eukaryotic cells, by engineering the cells to contain a selectable marker gene with a site for generating a double strand break within the intron of the
                                                                                                                                 The present invention relates to an assay for detecting chromosomal rearrangements (CR) in eukaryotic cells. The assay comprises engineering the cell to contain a selectable marker gene with a site for generating adouble strand break (ds break) within an intron of the cell; generating the ds break; detecting cells which repair the ds break; isolating DNA from the detected cells, and characterising the CR. The present sequence is a PCR primer for yeast URA3; which was used in the present invention to construct a URA3: actin intron::HO cut site yeast strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a collection of binding groups for a family of nucleic acide comprising members of relative high and relative low significance, where the binding groups are selected to be capable to identify, alone or in combination, essentially all members of the family of nucleic acids of relatively high significance. The collection of binding groups is useful for typing of nucleic acid in a clinical sample, by contacting the nucleic acid with the collection and determining whether one or more binding groups bound to the nucleic acid of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for determining or typing samples, has groups capable to identify essentially nucleic acids of relatively high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO:331.
                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV-1 related binding molecule oligonucleotide sequence SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 molecule, HIV-1; human immunodeficiency virus type 1; transcriptase; binding group; ss.
                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1; Length 19;
Pred. No. 1.6e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                Sequence 19 BP; 4 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Goudsmit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 87; 166pp; English
                                                                                                    Col 18; 16pp; English
                                                                                                                                                                                                                                                                                                                                                                                                      1050 GGAGTCCAACGCGTCCAT 1067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Collection of binding groups especially clinical samples, all members of the family of
                                                                                                                                                                                                                                                                                                                                  0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-JUL-2000; 2000EP-00202611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUL-2000; 2000EP-00202611
                                                                                                                                                                                                                                                                                                                                                                                                                                   GGAGTTCAATGCGTCCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Loukachov VV, Van Gemen B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                   16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-156696/21.
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 significance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-MAY-2002
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                                                                                                    Example 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Binding reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1536
                                                                                                                                                                                                                                                                                                                                                                     Matches
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               comprises at least a part of a member of relatively high significance of a family of nucleic acids. The collection of binding groups is useful for diagnosing the severity of a disease caused by a pathogen containing a member of a family of nucleic acids. ABL88779 to ABL89321 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymerase chain reaction amplification using combined ligated sequences as a template and enrichment primers directing the amplification towards
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Orientation-directed construction of a construct comprises performing
                                                                                                                                                                                                                     Gaps
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                                                                                       oligonucleotide sequences used in the exemplification of the present
sample. This method is useful for determining whether the sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       orientation enrichment reaction; GER; 8s.
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                                                                                                                                                                               Score 14.8; DB 1; Length 19;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
0.4%; Score 14.8; DB 1; Length 19;
Best Local Similarity 88.9%; Pred. No. 1.6e+03;
Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2038 PCR primer used to create constructs using OER method
                                                                                                                                                                                                                     Indels
                                                                                                                                           G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 5 A; 6 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 35; 60pp; English.
                                                                                                                                                                                                                                                       3609 CGTTCTGTACTGTTACTG 3626
                                                                                                                                                                                                                                                                                                                                                                                     AAD44897 standard; DNA; 19 BP.
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                                                                                                                                                                               0.4%;
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                                                                                                                                             Sequence 19 BP; 6 A; 4 C; 5
                                                                                                                                                                                                                                                                                            CGTCCAGTACTGTTACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                  Local Similarity 88.9
les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the desired orientation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-667004/71.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ben-Asouli Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-AUG-2002
                                                                                                            invention
                                                                                                                                                                                                                                                                                            18
                                                                                                                                                                                                                                                                                                                                                                                                                         AAD44897;
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                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                     Matches
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RESULT 1540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method for orientation-directed construction of a construct comprising at least two nucleic acid segments of interest. The method involves performing polymerase chain reaction (PCR) amplification reaction using the combined ligated sequences as template and specific enrichment primers directing the PCR amplification towards the desired orientation where each combined ligated sequence is amplified in a separate reaction creating a combined product having phosphorylated vectors specifically for orientation. Airected constructs or genetic vectors specifically for orientation. The present sequence is a DNA fragment used to illustrate the orientation enrichment reaction (ORR)
                                                                                                                                                                                                                                                                                                                                                                                               Orientation-directed construction of a construct comprises performing polymerase chain reaction amplification using combined ligated sequences as a template and enrichment primers directing the amplification towards
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                               construction; orientation enrichment reaction;
                                                                                                                                                       DNA fragment II used to illustrate the OBR method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 5 A; 6 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1;
Pred. No. 1.6e+03;
0; Mismatches 2;
       3651 CTTGCTTGCCTGCAGGGC 3668
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure, Fig 1, 60pp, English.
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                            18 criscarscerscassic 1
                                                                                     AAD44911 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                11-FEB-2002; 2002WO-IL000104.
                                                                                                                                                                                                                                                                                                       12-FEB-2001; 2001IL-00141392
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                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                            (GENE-) GENE BIO APPL LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                   desired orientation.
                                                                                                                                                                                                                                                                                                                                                    Ben-Asouli Y, Osman F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       method of the invention
                                                                                                                                                                              Orientation-directed
                                                                                                                                                                                           genetic vector; OER;
                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-667004/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                     WO200264774-A2
                                                                                                                                                                                                               Unidentified
                                                                                                                                 13-DEC-2002
                                                                                                          AAD44911;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Matches
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AAD44914 standard; DNA; 19

RESULT 1539

13-DEC-2002 (first entry)

AAD44914;

AAD44914 ID AAD4 XX XX AC AAD4 XX DT 13-D

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Orientation-directed construction of a construct comprises performing polymerase chain reaction amplification using combined ligated sequences as a template and enrichment primers directing the amplification towards
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
  to generate pcDNAPKRwt construct using the OER method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                              Orientation-directed construction; orientation enrichment reaction; genetic vector; OER; PCR; primer; ss.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human chromosome 1p36-35 PCR primer SEQ ID NO:2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        enrichment reaction (OER) method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 2 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Fig 3A; 60pp; English.
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                                                                                                                                                                                                                                                                  11-FEB-2002; 2002WO-IL000104.
                                                                                                                                                                                                                                                                                                                     12-FEB-2001; 2001IL-00141392.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTTGCATGCCTGCAGGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3651 CTTGCTTGCCTGCAGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL44958 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                   (GENE-) GENE BIO APPL LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the desired orientation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-667004/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
PCR primer #2,
                                                                                                                                                                   WO200264774-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP2001321190-A.
                                                                                                                                                                                                                                                                                                                                                                                                                    Ben-Asouli Y,
                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                    22-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2001.
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(GENO-)

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The invention relates to a method for screening compounds for their ability to modify the binding of G protein-coupled receptor protein GPR7 to its polypeptide, ligands and their amides, esters and salts, by measuring the binding in the presence and assence of the test compound. The method is used for prevention and treatment of cibophobia, anorexia, loss of appetite, excessive appetite and a broad range of obesity-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Screening compounds that modify the binding of G-protein coupled receptor GPR7 to its ligands for treatment of obesity and cibophobia.
                                                                                                                                                                                                                                         The invention describes an isolated and purified polypeptide, comprising an amino acid sequence encoded by exon 16 of the myotonic dystrophy protein kinase (DMPK) gene. The invention is used to detect presence or risk of myotonic dystrophy. Myocardial infarction or a condition associated with muscle damage or dysfunction. This sequence represents a mutant of the CTG repeat isolated from the 3' UTR of the novel Mytonic dystrophy protein kinase (DMPK) isoform gene that results in supression of splicing into the 3' splice site of the gene
                                                                    A new isoform of myotonic dystrophy protein kinase includes a sequence encoded by exon 16 of the gene and is useful to detect presence or risk of myotonic dystrophy, myocardial infarction or a condition associated with muscle damage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G protein-coupled receptor; GPR7; primer; 88; anorectic; cibophobia; anorexia; appetite loss; excessive appetite; obesity.related disorder; addipocyte malignancy; obesity; excessive insulin; blood volume change; thyroid disorder; paediatric obesity; upper body obesity; dietary obesity; cardiac obesity; whole body adipocyte disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 2 A; 5 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 214; 222pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Screening method related primer #40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3194 CCCCGGAGCTGGAGGATC 3211
                                                                                                                                                                                                    Example, Page 9; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-MAY-2002; 2002WO-JP004635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-2001; 2001JP-00145411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 88.9%;
tes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18 ccccadadcrccaddarc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (TAKE ). TAKEDA CHEM IND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABX92989 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mori M, Shimomura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-129320/12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                        The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. to array the multiwell conferration Nos. are mixed clones of the specified discrimination Nos second to discrimination Nos are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the calcultant cultures are amplified by using the above primer; (g) signals and laters are specified from the amplified products; (h) the clones are reconstituted as the positions on the chromosome and arrayed. The mixed clones are reconstituted as the positions on the chromosome and arrayed. The mixed represent PRR primers for human chromosome 1936-35 July, and ABL45323 to ABL45334 crepresent progressent invention continued in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 3 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                      Claim 4; Page 44; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1037 GACAGGTGTCCCTGGAGT 1054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GACAGGTGACCCTGGTGT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                       10-MAR-2000; 2000JP-00066716.
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                                                                      (RIKA ) RIKAGAKU KENKYUSHO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK87500 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 88.5 es 16; Conservative
                                                                                                                                                                                                    Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MAHA/) MAHADEVAN M S. (TISC/) TISCORNIA G.
                                                                                                                                                WPI; 2002-144136/19
                                                                                                 GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2002061571-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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RESULT 1541 ABK87500/c

Best Loca Matches

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Gaps

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ss; siNA; human; BCL2; short interfering nucleic acid; RNA interference; cytostatic; immunosuppressive; virucide; anti-HIV; cancer; autoimmune disease; viral infection; HIV.

WO2003070969-A2. Homo sapiens.

Human BCL2 siNA upper sequence SEQ ID NO:177.

(first entry)

12-FEB-2004

ADF49449;

19 GACGAGCTCTTCACGCTG

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ADF49449/c ID ADF49449 standard; RNA; 19

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disorders including adipocyte malignancy, obesity due to external factors, excessive insulin, blood volume changes, thyroid disorders, paediatric obesity, upper body obesity, dietary obesity, cardiac obssity and whole body adipocyte disorder. This sequence represents a primer used in the scope of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to novel compositions for inhibiting body weight gain, for lowering body weight, for inhibiting fat weight gain, and for suppressing appetite, which contain as active component a peptide ligand (GPRBL, ADC51805) of brain origin. The compositions can be used for treatment mastocytosis, exogenous obesity, hyperinsulinemic obesity, hyperplasmic obesity, hyperplasmic obesity, hyperplasmic obesity, hypothyseal obesity, hypothyseal obesity, hypothyseal obesity, symptomatic obesity, infant obesity, upper body obesity, alimentary obesity, hypogonadal obesity, systemic mastocytosis, simple obesity and central obesity). The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Composition comprising peptide of brain origin binding to orphan G-protein coupled receptor GPR8 for treatment and prevention of obesity and hyperphagia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                weight; GPR8L; brain; hyperphagia; obesity; anorectic; GPR8; PCR;
                                                                                                                                                                                Gaps
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                                                                                                                                           0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03; ive 0; Mismatches 2; Indels
                                                                                                        Sequence 19 BP; 4 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence 19 BP; 4 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matsumoto H, Noguchi J, Harada M, Mori M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 79; SEQ ID NO 148; 277pp; Japanese.
                                                                                                                                                                                                               1876 GAGGAGCTCTTCAAGCTG 1893
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-DEC-2001; 2001JP-00403260.
28-MAR-2002; 2002JP-00093096.
                                                                                                                                                                                                                                       19 GACGAGCTCTTCACGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                             GPR8 PCR primer, SEQ ID 148.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                                                                                                                                                                                      ADC51937 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                            Local Similarity 88.9
hes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Body weight primer; ss.
                                                                                                                                                                                                                                                                                                                                                                         ADC51937;
                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                   RESULT 1543
                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel short interfering nucleic acid (siNA) that downregulates expression of the BCL2 gene by RNA interference. A siNA of the invention has cytostatic, immunosupressive, virucide, and anti-HIV activity. The siNA are useful for modulation (inhibition) of expression or activity of BCL2 by RNA interference. siNA are used to modulate expression of BCL2 genes, in cells, tissue explants or organisms, e.g. for treating cancer, autoimmune diseases and viral infections (including by HIV) but also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (e.g. of sing represent siNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer or autoimmune disease, downregulates expression of the BCL2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 1 A; 5 C; 8 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 177; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1262 AGGACCGGCCGCCAAGC 1279
                                                                                                                                                                                                                                                                                   20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
18-UUL-2002; 2002US-0396905P.
29-AUC-2002; 2002US-0406784P.
05-SEP-2002; 2002US-040878P.
                                                                                                                                                                                                                                                        18-FEB-2003; 2003WO-US004908.
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                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 88.9
les 16; Conservative
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                                                                                                                                                                                                                           28-AUG-2003
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Matches
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Gaps

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Query Match 0.4%; Score 14.8; DB 1; Length 19; Best Local Similarity 88.9%; Pred. No. 1.6e+03; Matches 16; Conservative 0; Mismatches 2; Indels

1876 GAGGAGCTCTTCAAGCTG 1893

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The invention relates to a novel short interfering nucleic acid (siNA) that downregulates expression of the BCL2 gene by RNA interference. A siNA of the invention has cytostatic, immunosuppressive, virucide, and anti-HIV activity. The siNA are useful for modulation (inhibition) of expression or activity of BCL2 by RNA interference. siNA are used to modulate expression of BCL2 genes, in cells, tissue explants or organisms, e.g. for treating cancer, autoimmune diseases and viral infections (including by HIV) but also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (e.g. of single runclectide polymorphisms). The sequences shown in ADF49273-ADF50143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                 ss; siNA; human; BCL2; short interfering nucleic acid; RNA interference; cytostatic; immunosuppressive; virucide; anti-HIV; cancer; autoimmune disease; viral infection; HIV.
                                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer or autoimmune disease, downregulates expression of the BCL2 gene.
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18.9%; Pred. No. 1.6e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 5 A; 8 C; 5 G; 0 T; 1 U; 0 Other;
                                             Human BCL2 siNA upper sequence SEQ ID NO:591
                                                                                                                                                                                                                                                                                                                                                                             Example 3; SEQ ID NO 591; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1262 AGGACCGGCCGCCAAGC 1279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGGACCAGGCCUCCAAGC 19
                                                                                                                                                                                              2002US-0363124P.

2002US-0386482P.

2002US-0386905P.

2002US-0406784P.

2002US-0408378P.

2002US-0409233P.
                                                                                                                                                                  18-FEB-2003; 2003WO-US004908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.9%;
                                                                                                                                                                                                                                                                             RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF31538 standard; RNA; 19
                                                                                                                                                                                                                                                                                                 Beigelman L;
                          12-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                   WPI; 2003-712622/67.
                                                                                                                           #02003070969-A2
                                                                                                                                                                                                                  18-JUL-2002;
29-AUG-2002;
                                                                                                                                                                                               11-MAR-2002;
                                                                                                                                                                                                                                      05-SEP-2002;
                                                                                                                                                                                                                                                         15-JAN-2003;
                                                                                                                                                                                                                                                                                                 Mcswiggen J,
                                                                                                         Homo sapiens
                                                                                                                                               28-AUG-2003
         ADF49863;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADF31538
1D ADF3
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AC ADF3
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DE Huma
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human insulin-like growth factor 1

comprise ribonucleotides and may be double or single stranded. They comprise sense and antisenes regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the IGF-IR gene in cells, tissue explants for the captering of an ariety of conditions. They may be used for treating cancer and other proliferative diseases (e.g., restension and polycystic kindney disease), inflammatory and/or allergic diseases, autoimmune diseases and transplant rejection. The siNAs are also useful for drug sorreining, diagnosis, therapeutic target identification and validation, sorreining, diagnosis, therapeutic target identification and validation, sorreining, diagnosis, therapeutic target identification and validation, and contains the sinas and some and capted contains and capted capted identification and validation, and contains the sinas are also asful for drug sorreining, diagnosis, therapeutic target identification and validation, and contains the sinas and succession and validation, and contains the sinas and succession and admits and contains the sinas and successions and admits and contains the sinas and successions and admits and contains the sinas and successions and admits and contains and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human IGF-1R-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                 miRNA:
RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA, double-stranded RNA; micro-RNA; miRNA short hairpin RNA; siRNA; double-stranded RNA; micro-RNA; miRNA short hairpin RNA; shRNA; axpression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; proliferative disease; allergic disease; aucimmune disease; transplant rejection; cytostatic; vasotropic; nephrotropic; transplant rejection; cytostatic; immunosuppressive; human; antiinflammatory; autiallergic; immunosuppressive; human; insulin-like growth factor 1 receptor; IGF-IR; target sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    siNA, which is identical to the IGF-1R transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 203; 147pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002US-0363124P.
2002US-0386782P.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-FEB-2003; 2003WO-US005044
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 66.7
les 12, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-AUG-2003
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Matches
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1816 GGGGTCCTGCTCTGGGAG 1833

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Human IGF-1R transcript target sequence/siNA upper strand, SEQ ID NO:203.

(first entry)

12-FEB-2004

ADF31538;

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mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human IGF-IR-targeted double-stranded siNA.

Sequence 19 BP; 4 A; 9 C; 5 G; 0 T; 1 U; 0 Other;

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short interfering nucleic acid, useful e.g. for treatment and mosis of cancer , downregulates expression of the insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; micro-RNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; short interfering interfering size interfering; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; aucoimmune disease; transplant rejection; cytostatic; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human; insulin-like growth factor I receptor; IGF-IR; ss.
                                                                                                                                                                                                                                                                                                                                                                                        Human IGF-1R siNA lower strand, SEQ ID NO:480.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chowrira B;
2 GGGGUCGUCCUCUGGGAG 19
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11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
                                                                                                                                                                             ADF31815 standard; RNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                     (first entry)
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factor-l receptor gene.
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                                                                                                                                                                                                                                                                                                                     12-FEB-2004
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                                                                                                                                                                                                                                               ADF31815;
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The present invention describes a method for enhancing the expression of a protein of interest from Bacillus. The method comprises: (a) obtaining an altered Bacillus strain capable of producing a protein of interest, where the altered Bacillus strain has at least one inactivated chromosomal gene consisting of sbo, sir, ybc0, csn, spollSA, sigB, phrC, rapA, CssS, trpA, trpB, trpC, trpD, trpE, trpF, tdh/Kbl, alsD, sigD, prC, gapB, pckA, fbp, rocA, ycgN, ycgM, rocF, and rocD; and (b) growing the altered Bacillus strain under conditions such that the protein of interest is enhanced compared to the expression of the protein of interest is enhanced compared to the expression of the protein of interest is enhanced compared to the expression of the protein of interest is enhanced compared to the expression of the protein of interest is enhanced compared to the expression of the protein of interest in an unaltered Bacillus host strain. Also described: (1) an altered Bacillus strain comprising a chrow sigD, prpC, gapB, pckA, fbp, rocA, ycgN, ycgN, rocF, and conf. (csn, spollSA, sigB, phrC, rapA, CssS, trpA, trpB, trpC, trpD, trpE, trpF, trbP, alsD, sigD, prpC, gapB, pckA, fbp, rocA, ycgN, ycgN, rocF, and the DNA construct; (4) a host cell comprising the plasmid; (5) a method for obtaining an altered Bacillus strain with enhanced protease in an expediction; (6) a method for enhancing expression of a protease in an
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                                                                                                                                                                                                                                                                                                                                                                                                      Bacillus; sbo; slr; ybcO; csn; spollSA; sigB; phrC; rapA; CssS; trpA; trpB; trpC; trpD; trpE; trpF; tdh/kbl; alsD; sigD; prpC; gapB; pckA; fbp; rocA; ycgN; ycgM; rocF; rocD; enhancing expression; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Enhancing expression of a protein of interest from Bacillus by obtaining an altered Bacillus strain capable of producing a protein of interest and growing the altered Bacillus strain.
                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                Bacillus subtilis sigD deletion construct PCR primer SEQ ID NO:186
                                                ;
0
    Score 14.8; DB 1; Length 19;
Pred. No. 1.6e+03;
                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ferrari E, Harbison C, Rashid MH, Weyler W;
                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 186; 114pp; English.
                                                                                        1816 GGGGTCCTGCTCTGGGAG 1833
                                                                                                                       18 GGGGTCGTCCTCTGGGAG 1
                                                                                                                                                                                                                                       BP
0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAR-2002; 2002US-0368858P.
29-MAR-2002; 2002US-0368949P.
29-APR-2002; 2002US-0376343P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-MAR-2003; 2003WO-US009585.
                                                                                                                                                                                                                                       ADF41858 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                      12-FEB-2004 (first entry)
                      Local Similarity 88.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GEMV ) GENENCOR INT INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-876987/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
Bacillus subtilis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003083125-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-OCT-2003
                                                                                                                                                                                                                                                                               ADF41858;
      Query Match
                                                                                                                                                                                             RESULT 1548
                                              Matches
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human insulin-like growth factor 1 creeptor (IGF-IR) gene by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They comprise sense and may be double or single stranded. They comprise sense and may be double or single stranded. They comprise sense and antisense regions, or alternatively are comprise sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits of the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the IGF-IR gene in cells, tissue explants or organisms of the IGF-IR gene in cells, tissue explants or organisms or treatment of a variety of conditions. They may be used for treating cancer and other proliferative diseases (e.g., restenosis and polycystic diseases) inflammatory and/or allergic diseases, autoimmune diseases), inflammatory and/or allergic diseases, autoimmune chiseases), inflammatory and/or allergic diseases, autoimmune chiseases, inflammatory and/or allergic diseases and transplant rejection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene

Example 3; SEQ ID NO 480; 147pp; English.

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altered Bacillus; (7) a method for enhancing the expression of a protein of interest in Bacillus; (8) a method for enhancing the expression of a protein of interest in Bacillus; (9) a method for obtaining a protein of interest from a Bacillus; (10) an isolated nucleic acid comprising a sequence having e.g., 721, 857 or 995 bp and encoding a sequence having e.g., 211, 387 or 433 amino acids; and (11) an isolated amino acid sequence comprising e.g., 211, 387 or 433 amino acids are the method is useful for enhancing expression of a protein of interest from Bacillus. The present sequence is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-v-abl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; BCR; ss; siRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel double-stranded short interfering
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human breakpoint cluster region-targeted siRNA - SEQ ID 273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1; Length 19;
Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 6 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3742 ATTITGTTAACCCAGCGA 3759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-MAR-2002; 2002US-0363124P.

06-JUN-2002; 2002US-03867B2P.

15-AUG-2002; 2002US-0404039P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

05-SEP-2002; 2002US-0409378P.

14-JAN-2003; 2003US-040923P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF83979 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 88.9
es 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-679889/64.
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                                                                                                                                                                                                                                                                                                                               invention.
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ADF83979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of leukemia and lymphoma, downregulates the breakpoint cluster region-Abelson (BCR-ABL) gene.
                                                                                                             Gaps
 The current sequence is that
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                                                                                                                                                                                                                                                                                                                                                                                  short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; BCR; ss; siRNA.
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                                                                                                                                                                                                                                                                                                                                                      Human breakpoint cluster region-targeted siRNA - SEQ ID 10.
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88.9%; Pred. No. 1.6e+03;
                                                                              Score 14.8; DB 1; Length 19;
Pred. No. 1.6e+03;
                                                                                                             Indels
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                                              Sequence 19 BP; 1 A; 9 C; 7 G; 0 T; 2 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 2 A; 7 C; 9 G; 0 T; 1 U; 0 Other;
gene function studies and gene mapping. The cuthe human BCR-targeted siRNA of the invention.
                                                                                                            1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J, Beigelman L, Chowrira B;
                                                                                                                                             231 CTGGACACGGCCCGAGCG 248
                                                                                                                                                               2 CUGGCCACGCCCGCGCG 19
                                                                                                                                                                                                                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
15-AUG-2002; 2002US-0404039P.
29-AUG-2002; 2002US-0406784P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
14-JAN-2003; 2003US-0439922P.
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                                                                              0.48;
                                                                                              83.3%;
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                                                                                                             Matches 15; Conservative
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Matches 16; Conserv
                                                                                              Best Local Similarity
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2943 AGGGAGGCCCCAGGGGTC 2960
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2002US-0386782P.
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2002US-0408378P.
2002US-0409293P.
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2003US-0440129P.
                                                                                                                                                                          83.3%;
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                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                         Query Match
Best Local Similarity 83.3
Matches 15; Conservative
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06-JUN-2002;
29-AUG-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                    ADG34745;
                                                                                                                                                                                                                                                                                            RESULT 1552
                                                                                                                                                                                                                                                                                                                          88888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to short interfering nucleic acids (siNA) which downrequlate expression of the human tumour necrosis factor (TWF) gene by MA interference. The siNas may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense conjectucle and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmedified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised, expressed from a vector of sink, and vectors that express sink. The sinks are used to modulate expression of the TNF gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafte and transplants for the treatment of a variety of conditions. The TNF sinks have antibacterial, immunosuppressive, anticheumatic, antiatrhritic, anti-HIV, antipsoriatic and
                                                                                                                                                                                                                          RNA interference; short interfering nucleic acid; siNA; short siRNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shoresion modulation; gene therapy; dragnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; tumour necrosis factor; TNF; human; DNA-RNA hybrid; sa, antibacterial; immunosuppressive; antitheumatic; antiarthritic; anti-HIV; antipsoriatic; antiinflammatory; septic shock; rheumatoid arthritis; HIV/AIDS; psoriasis; inflammation; autoimmune disease; target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid, useful e.g. for treatment and diagnosis of septic shock or rheumatoid arthritis, downregulates expression of the tumor necrosis factor gene.
                                                                                                                                                                                                   Human TNF siNA oligonucleotide SEQ ID NO:9.
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                                    18 crescencecceced 1
                                                                                                             ADG34657 standard; RNA; 19 BP
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2002US-0386782P.
2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-SEP-2002; 2002US-0408378P 09-SEP-2002; 2002US-0409293P
       231 CTGGACACGGCCCGAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-0429359P
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                                                                                                                                                                      (first entry)
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28-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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continuing lamancoty activities. They way be used for treating especies shock, cheeses. The sinks are also useful for drug screening, dispensis, cheeses. The sinks are also useful for drug screening, dispensis, cheeses. The sinks are also useful for drug screening, dispensis, cheeses. The sinks are also useful for drug screening, dispensis, cheeses. The sinks are also useful for drug screening, dispensis, cheeses. The sinks are also useful for drug screening, dispensis to cheese the service of the service and the service
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(MCSW/) MCSWIGGEN J A. (BEIG/) BEIGELMAN L.

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may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in synthesised. The invention also relates to kits for the in vitro or in synthesised. The invention also relates to kits for the in vitro or in that express siNA. The siNAs are used to modulate expression of the TNF gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. The TNE siNAs have antibacterial, immunosuppressive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Virucide; Hepatotropic; Gene therapy; 8s; short interfering nucleic acid; siNA; hepatitis B virus; HBV; RNA interference.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    antifheumatic, antiathritic, anti-HIV, antipsoriatic and antitheumatic, antiathritic, anti-HIV, antipsoriatic and antiinflammatory activities. They may be used for treating septic shock, rheumatoid arthritis, HIV/AIDS, psoriasis, inflammation and autoimmune diseases. The siMhs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TNP-targeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 1 A; 9 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
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20-MAR-2000; 2000US-00436430.
24-OCT-2000; 2000US-00631025.
08-JUN-2001; 2001US-00874778.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
20-FEB-2002; 2002US-0358580P.
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94US-00193627
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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ADM00745/C
XX
AC ADM00745
AC ADM00745
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DT 20-MAY-2
XX
XX
BE Hepatiti
XX
XX
BE 14-MAY-1
BE 09-MAR-2
BE 09-M
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   8
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2002US-0409293P

(MORR/) MORRISSEY D

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The invention relates to a short interfering nucleic acid (sinh) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA interference or that inhibite HBV replication. Also disclosed are the following: (i) a method of modulating the expression of a HBV gene in a following: (i) a method of generating a library of sinh constructs tissue explant; (ii)) a method of generating a library of sinh constructs complexity; (iii) a cell containing one or more sinh molecules; (iv) a kit containing a sinh molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or computate the expression of a HBV target gene in a cell, tissue or molecule is adapted for use to treat HBV infection, and comprises sequence complementary to an RNA sequence encoding HBV and the sense region comprises sequence complementary to the antisense region comprises the sense region and the sense region of the sinh molecule is assembled from 2 nuclecule, where sense region mad the confisense region of the sinh molecule. Where sense region and the sense region and the sense region of the sinh molecule. The linker molecule is a polynucleotide inker molecule. The linker molecule is a polynucleotide confisense region of the antisense region of the sinh all the sense region comprises a 3 - terminal overhang and the antisense region comprises a 3 - terminal coverhang and the antisense region comprises a 3 - terminal coverhang and the antisense region comprises a 3 - terminal coverhang and the antisense region comprises a conformentary to RNA cencoding HBV. The siNA is useful for treating human hepatitis B virus infections, and for characterising pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function conformation between in genes function analysis. The sinh molecules in spender represents 1 of 1504 HBV with a sunfor research settings.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Virucide; Hepatotropic; Gene therapy; ss; short interfering nucleic acid; siNA; hepatitis B virus; HBV; RNA interference.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                 New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.8; DB 1; Length 19; 88.9%; Pred. No. 1.6e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 8 A; 2 C; 8 G; 0 T; 1 U; 0 Other;
                                                             Morrissey D, Mcswiggen JA, Beigelman L;
                                                                                                                                                                                                                                                                      Claim 11; Page 48; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sinA molecules of the invention.
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                                                                                                                                                                                                                        characterizing gene function.
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Matches 16; Conservative
                                                                                                        WPI; 2003-901032/82.
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1 UUCCUCUUCAUCCUGCUG 18

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RESULT

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Claim 11; Page 48; 72pp; English.
                         07-FEB-1994; 94US-00193627.
08-NOV-1999; 99US-00436430.
20-MAR-2000; 2000US-00531025.
09-AUG-2000; 2000US-00636385.
24-OCT-2000; 2000US-00696347.
                                                           08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
                                                                                      20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
26-MAR-2002; 2002WO-US009187.
                                                                               05-DEC-2001; 2001US-0337055P
                                                                                                           06-JUN-2002; 2002US-0386782P.
                                                                                                                  2002US-0406784P
2002US-0408378P
     16-SEP-2002; 2002US-00244647
                                                                                                                               2002US-0409293P
                                                                                                                                                                         Mcswiggen JA,
                                                                                                                                            (MORR/) MORRISSEY D.
(MCSW/) MCSWIGGEN J.A.
(BEIG/) BEIGELMAN L.
                                                                                                                                                                                      WPI; 2003-901032/82.
                                                                                                                                                                        Morrissey D,
                                                                                                                               09-SEP-2002;
                                                                                                                  29-AUG-2002;
05-SEP-2002;
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The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA interference or that inhibits HBV replication. Also disclosed are the collowing: (i) a method of modulating the expression of a HBV gene in a tissue explant; (ii)) a method of generating a library of siNA constructs molecules; (iv)a kit containing a siNA molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or modulate the expression of a HBV target gene in a cell, tissue or conjulate the expression of a HBV target gene in a cell, tissue or molecule is adapted for use to treat HBV infection, and comprises a sense of molecule is adapted for use to treat HBV infection, and comprises sequence complementary to an RNA sequence encoding HBV and the sense region comprises sequence complementary to the antisense region. Where the silva comprises the sense region and the second fragment comprises the comprises the sense region and the second fragment comprises the antisense region of the silva molecule. Where sense region and the cantisense region of the silva molecule is a polynucleotide. Comprises separate oligonucleotides, and are covalently connected via a linker molecule. The linker molecule is a polynucleotide. The sense region comprises a 3'-terminal overhang and the antisense region 3'-terminal overhang is complementary to RNA encoding HBV. The silva is useful for treating human hepatitis B virus concing HBV. The silva is useful for treating human hepatitis B virus concing hBV. The silva is useful for treating human hepatitis B virus confidence of unkibit activity of target genes in a pathway to determine the function of unkibit activity of target genes in a pathway to determine the function may also he used in clinical, industrial, enhancemental, and per function may also he used in clinical, industrial, he with may also he used in a plant function may also he used in clinical industrial industrial industrial industrial industria
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 siNA molecules of the invention.
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Human HER2 (EGFR2) transcript target sequence/siNA upper strand, SEQ:68.
                                                                                                   RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analyysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; erbB2; c-erb-B-2; target sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fosnaugh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 68; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J, Pavco P, Beigelman L,
BP
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09-SEP-2002; 2002US-0409293P.
19-SEP-2002; 2002US-00251117.
21-OCT-2002; 2002US-0027494.
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2002US-00163552.
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03-JUL-2002; 2002US-0393924P.
29-AUG-2002; 2002US-0406784P.
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ADL78903 standard; RNA; 19
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                                                   20-MAY-2004 (first entry)
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                                                                                                                                                                                                                                       WO2003070912-A2.
                                                                                                                                                                                                              Homo sapiens.
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06-JUN-2002;
                                                                                                                                                                                                                                                                                                                    20-FEB-2002;
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                        ADL78903;
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                                                                                                                                                                                                                                                                                                  New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for characterizing gene function.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 14.8; DB 1; Length 19;
14.4%; Pred. No. 1.6e+03;
ive 8; Mismatches 2; Indels
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                                                                                                                                                                                                                                                     Beigelman L;
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Best Local Similarity 44.4
Matches 8; Conservative
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Jamison S;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth factor receptor gene.
                     studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human HER2 (EGFR2)-targeted double-stranded siNA, which is identical
identification and validation, genetic engineering, pharmacogenomics,
                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA interference, short interfering nucleic acid; siNA, short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; erbB2; c-erb-B-2; target sequence; ss.
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                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 19; 72.2%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                            2; Indels
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                                                                                                                                Sequence 19 BP; 3 A; 5 C; 7 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                            3; Mismatches
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                                                                                          the HER2 transcript target sequence.
                                                                                                                                                                                                                                                                          1582 CGGGCCATGGAGTACTTG 1599
                                                                                                                                                                                                                                                                                                    CUGGGCAUGGAGCACUUG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL79015 standard; RNA; 19 BP
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11-MAR-2002; 2002US-0363124P.
29-MAY-2002; 2002WO-US016840.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
19-SEP-2002; 2002US-00251117.
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06-JUN-2002; 2002US-0386782P
03-JUL-2002; 2002US-0393924P
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                                                                                                                                                                                                                            13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-697612/66.
                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                               Query Match
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ADL/79015
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AC ADL/79011
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BNA inct
KW Short inc
KW Short
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an antisense oligonucleotide. Specifically, the sinks include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The sinks can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, capressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sink, conjugates and/or complexes of sink, and vectors that express sink. Conjugates used to modulate expression of EGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The sinks are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide pluman HER2 (EGFR2)-targeted double-stranded sink, which is identical to
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or alternatively are assembled from a sense oligonucleotide and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 19; 72.2%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human HER2 (EGFR2) sinA lower strand, SEQ ID NO:317.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 3 A; 2 C; 9 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the HER2 transcript target sequence.
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2002US-00163552.
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2002US-0393924P.
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09-SEP-2002; 2002US-0409293P.
19-SEP-2002; 2002US-00251117.
21-OCT-2002; 2002US-0027494.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              72.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADL79152 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 72.2
les 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003070912-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADL79152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1557
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Matches
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dwortegulate expression of one or more human epidermal growth factor receptor (BGFR) genes (including HER1, HER2 HER3 and HER4) by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and mantisense regions, or alternatively are assembled from a sense oligonuclectide and mitsense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and confund modified expression of EGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment of a variety of conditions, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a HERZ (EGFRZ)-targeted double-stranded sinA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                          New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                              The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA interference; short interfering nucleic acid; siNA; short interfering many siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; BGFR; HER2; EGFR2; neu; exbB2; c-erb-B-2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 19;
88.9%; Pred. No. 1.6e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human HER2 (EGFR2) sinA lower strand, SEQ ID NO:429.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 7 C; 5 G; 0 T; 3 U; 0 Other;
                                                                                                                                 Example 3; SEQ ID NO 317; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1582 CGGGCATGGAGTACTTG 1599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL79264 standard; RNA; 19 BP
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29-MAY-2002; 2002WO-US016840.
06-UUN-2002; 2002US-00163552.
66-UUN-2002; 2002US-0366782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2003; 2003WO-US005045
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match
Best Local Similarity 88.>*,
Best Local Similarity 16,
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAY-2004 (first entry)
                                                                                           factor receptor gene.
WPI; 2003-697612/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003070912-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1558
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor receptor (ESPR) genes (including HER1, HER2 and HER4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vector or enzymatically synthesised. The siNAs are used to modulate expression of EGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants the treating a wide range of cancers such as breast and ovarian cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a betage of the mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic; vasotropic; nephrotropic; cerebroprotective; treating leukaemia; solid tumors; restenosis; polycystic kidney disease; bronchiolitis; glomerulonephritis; stroke; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short harripin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics;
                                                                                                                                                                                                                                                                                                                   New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1; Length 19; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PDGFr-targeted sinA upper strand SEQ ID NO:168.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene function analysis; gene mapping; human;
platelet derived growth factor receptor; PDGFr; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 5 A; 9 C; 2 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 429; 171pp; English.
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                29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-040223P.
19-SEP-2002; 2002US-00251117.
21-OCT-2002; 2002US-0027494.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match

Best Local Similarity 88.9%;
Matches 16; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19 CTGGTGGATGCTGAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              862 CTGGTGGAGGCTGACGAG
                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO14737 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                   factor receptor gene.
                                                                                                                                                                                                                                                                        WPI; 2003-697612/66.
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0.4%; Score 14.8; DB 1; Length 19; 61.1%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                            5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                          1801 GACGTCTGGTCCTTTGGG 1818
                                                                                                                                                                                                                                                                                                                                                                                                 |||||::||:|||
GACGUGUGGUCCUUCGGG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                           05-FEB-2003; 2003WO-US003473
                                                        2002US-0358580P
                                                              2002US-0363124P
2002US-0386782P
                                                                           2002US-0406784P
                                                                                 2002US-0408378P
                                                                                        2002US-0409293P
                                                                                             15-JAN-2003; 2003US-0440129P
                                                                                                                       Mcswiggen J, Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO15048 standard; RNA; 19
                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                             11; Conservative
                                                                                                                                    WPI; 2003-731605/69.
                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                  WO2003072704-A2
       Homo sapiens.
                                                              11-MAR-2002;
06-JUN-2002;
                                                                          29-AUG-2002;
05-SEP-2002;
                                                                                        09-SEP-2002;
                               04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADO15048;
                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1560
ADO15048/c
                                                                                                                                                                                                                                                                                                                                                                             Matches
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cytostatic; vasotropic; nephrotropic; cerebroprotective; treating leukaemia; solid tumors; restenosis; polycystic kidney disease; bronchiolitis; glomerulonephritis; stroke; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; New short interfering nucleic acid, useful e.g. for treatment and diagnosis of tumors, downregulates expression of the platelet-derived growth factor receptor gene. expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; human; platelet derived growth factor receptor; PDGFr; 88. Human PDGFr-targeted sinA lower strand SEQ ID NO:479. Example 3; SEQ ID NO 479; 148pp; English 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. 05-FEB-2003; 2003WO-US003473 Mcswiggen J, Beigelman L, (RIBO-) RIBOZYME PHARM INC WPI; 2003-731605/69. WO2003072704-A2 Homo sapiens. 20-FEB-2002; 11-MAR-2002; 36-JUN-2002; 04-SEP-2003. The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human platelet-derived growth factor ceceptor (PDGFT) gene by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They turther comprise sense and may be double or single stranded. They curther comprise sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA, doublestranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA, doublestranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain also relates to kits complexes and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siRNA; conjugates and/or complexes of siRNA; and vectors that express siNA. The siNAs are used to modulate expression of the PDGFr gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating leukaemia and solid tumours, restenosis, polycystic kidney clasease, bronchiolitis, glomerulonebritis and stroke. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification condition, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human PDGFr transcript targeted doubble-stranded siNA, which is identical to the PDGFr transcript New short interfering nucleic acid, useful e.g. for treatment and diagnosis of tumors, downregulates expression of the platelet-derived growth factor receptor gene. Seguence 19 BP; 2 A; 4 C; 8 G; 0 T; 5 U; 0 Other; Example 3; SEQ ID NO 168; 148pp; English. Chowrira B;

Chowrira B;

2002US-0363124P. 2002US-0386782P.

2002US-0358580P

The invention relates to short interfering nucleic acids (SINA) which downregulate expression of the human platelet-derived growth factor downregulate expression of the human platelet-derived growth factor ceceptor (PDGFT) gene by RNA interference. The SINAs may or may not comprise ribonucleotides and maisenes regions, or alternatively are further comprise sense and antisenes regions, or alternatively are sasembled from a sense oligonucleotide and a natisense oligonucleotide. Specifically, micro-RNA (miRNA) and short interfering RNA (siRNA, double-certain burnmodified or chemically modified, can contain also relates to kits deoxyribonucleotides, and can be chemically synthesised, expressed from a corpus and vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siRNA, conjugates and/or complexes of siRNA; and vectors that express siNA. The siNAs are used to modulate expression of the PDGFT gene in cells, tissue explants or corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants to the treatment of a variety of conditions. They may be used for treating leukaemia and solid tumours, restences; polycystic kidney disease, bronchiolitis, glomerulonephritis and stroke. The siNAs are also and validation, canaetic endiaperion charmanchemic a stroke are also and validation, canaetic endiaperion charmanchemic a stroke. ö function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human PDGFr-targeted double-stranded siNA, which is identical to the PDGFr transcript and validation, genetic engineering, pharmacogenomics, studying gene unction, and gene mapping (e.g., of single nucleotide polymorphisms) the present sequence represents the lower strand of a human PDGFr-Gaps ö Score 14.8; DB 1; Length 19; Pred. No. 1.6e+03; 0; Mismatches 2; Indels Sequence 19 BP; 5 A; 8 C; 4 G; 0 T; 2 U; 0 Other; 0.4%; Best Local Similarity 88.9 Matches 16; Conservative and validation, target sequence Query Match

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Gaps

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2; Indels

1801 GACGTCTGGTCCTTTGGG 1818

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ABD24824

ABD24824

XXX
XXX
ABD3

XXX
ABD3

XXX
ABD4

XXX
ABD7

AB 음

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19 GACGTGTGGTCCTTCGGG
                                                                                                                               ABD24824 standard; DNA; 19
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ABD24824;

(first entry) 29-JUL-2004 AI092623-derived oligonucleotide SEQ ID 3836.

analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;

Homo sapiens

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

ä Aguilar Pabalan J, Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, 1 Miller S,

WPI; 2003-093058/08.

to Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent

Claim 15; SEQ ID NO 3836; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating branchoconstriction, respiratory tract inflammation, allergies and reducing adenosine ensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic. with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to pulmonary obstruction, and/or bronchoconstriction and/or lungs inflammation, allergies and/or surfactant hypoproduction are associated

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                                                                                                                                                                                                                        ss; detection; RNA interference; siRNA; gene silencing; gene expression;
the oligonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                       Gaps
                                                                      ;
0
                                                      Length 19;
                                                                      Indels
                                                                      7
                                    BP; 3 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                      DB 1;
                                                   Score 14.8; DB 1;
Pred. No. 1.6e+03;
0; Mismatches 2
                                                                                                                                                                                                      RNA interference target sequence #34.
                                                                                       2979 GACCAGGGCTTTTTCTGG 2996
                                                                                                        1 GACCAAGGCTTTTCCTGG 18
                                                                                                                                                     ВР.
                                                                                                                                                                                                                                                                                                     21-NOV-2003; 2003WO-JP014893.
                                                                                                                                                                                                                                                                                                                     22-NOV-2002; 2002JP-00340053
                                                              88.9%;
                                                                                                                                                   ADQ27126 standard; DNA; 19
                                                                                                                                                                                      (first entry)
                                                    Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                     (NATO/) NATORI Y.
(SAIG/) SAIGO K.
(TEIK/) TEI K.
(NAIT/) NAITO Y.
                                                                                                                                                                                                                                                                  WO2004048566-A1.
                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                 cytotoxicity
                                   Sequence 19
                                                                                                                                                                                      26-AUG-2004
                                                                                                                                                                                                                                                                                   10-JUN-2004.
                                                                                                                                                                    ADQ27126;
                                                                                                                                  RESULT 1562
                                                                                                                                            ADQ27126
  SSSSS
                                                                                        8
                                                                                                        셤
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.. 0 The invention relates to a method of detecting the base sequence for RNA interference by detecting the regions in the DNA sequence fulfilling the following requirements such as: (i) the base at 3' terminal is adenine, thymine or uracil; (ii) the base at 5' terminal is guanine or cyclosine; (iii) the seven base sequence at 3' terminal is guanine or cyclosine; and uracil, and; (iv) there are bases in a such a number that it causes RNA interference without showing cyclotoxicity. The method is used for designing and synthesizing siRNA causing RNA interference. This sequence corresponds to an RNA interference target sequence of the invention. Gaps ; 0 Score 14.8; DB 1; Length 19; Pred. No. 1.6e+03; 0; Mismatches 2; Indels Sequence 19 BP; 5 A; 3 C; 7 G; 4 T; 0 U; 0 Other; 594 18 0.4%; 577 GGCAGCGACGTGGAGTTC 1 GGTAGCAACGTGGAGTTC Query Match 0.4 Best Local Similarity 88.9 Matches 16; Conservative 8 셤

Detecting sequence of RNA interference useful for synthesizing siRNA, by detecting regions in sequence fulfilling specific criteria such as base at 3' terminal is adenine, thymine or uracil, base at 5' terminal is guanine or cytosine.

Naito Y;

Tei K,

Saigo K,

WPI; 2004-487423/46.

Disclosure; SEQ ID NO 48; 325pp; Japanese.

Ι

RESULT 1563

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This invention relates to a novel method of introducing a nucleic acid into an adipocyte which comprises contacting an adipocyte having a cell membrane with a nucleic acid molecule, thus forming a mixture and electroporating the mixture under conditions such that the cell membrane becomes permeabilised, such that the nucleic acid is introduced into the adpocyte. The invention may be useful for the production of compounds with an antidiabetic or anorectic activity whilst the disclosed sequences may be useful for gene therapy. The methods are useful for treating type if diabetes, insulin resistance or obesity. The present sequence is that of a region of a gene which may be targeted by short inhibitory RNA (SIRNA) used with the method of the invention.
permeabilised cell membrane; antidiabetic; anorectic; gene therapy;
type II diabetes; insulin resistance; obesity; Myolc; ds; mouse; murine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Introducing a nucleic acid into an adipocyte, useful for treating type diabetes, obesity or insulin resistance, comprises contacting an adipocyte having a cell membrane with a nucleic acid molecule, thus forming a mixture.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 4 A; 2 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 102; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%;
                                                                                                                                                                                                                     11-DEC-2003; 2003WO-US039774
                                                                                                                                                                                                                                                                     11-DEC-2002; 2002US-0432427P
                                                                                                                                                                                                                                                                                                                                                                       Jiang Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CAGGAGGATTGGATTTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT89883 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                     (UYMA-) UNIV MASSACHUSETTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-468860/44.
                                                                                                                                                                                                                                                                                                                                                                  Czech MP, Zhou Q,
                                                                                                                      WO2004053103-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ribonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DE19601385-A1
                                                                           Mus musculus.
                                                                                                                                                                      24-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT89883;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT89883
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of detecting the base sequence for RNA interference by detecting the regions in the DNA sequence fulfilling the following requirements such as: (i) the base at 3 'terminal is adenine, thymine or uracil; (ii) the base at 5' terminal is guanine or cytosine; (iii) the seven base sequence at 3' terminal is graine or cytosine; (iii) the seven base sequence at 3' terminal is rich in adenine, thymine and uracil, and; (iv) there are bases in a such a number that it causes RNA interference without showing cytotoxicity. The method is used for designing and synthesizing siRNA causing RNA interference. This sequence corresponds to an RNA interference target sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting sequence of RNA interference useful for synthesizing siRNA, by detecting regions in sequence fulfilling specific criteria such as base at 3' terminal is adenine, thymine or uracil, base at 5' terminal is
                                                                                                                                                                                                                                                                                            вв; detection; RNA interference; віRNA; gene silencing; gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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0.4%; Score 14.8; DB 1; Length 19;
Best Local Similarity 88.9%; Pred. No. 1.6e+03;
Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 6 A; 1 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 50; 325pp; Japanese.
                                                                                                                                                                                                                                               RNA interference target sequence #36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       651 GGTGAATGGCAGCAAGGT 668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
                                                                                                  ADQ27128 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-NOV-2003; 2003WO-JP014893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-2002; 2002JP-00340053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP48862 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Saigo K, Tei K, Naito Y;
                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              guanine or cytosine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-487423/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NATORI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SAIGO K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TEI K.
NAITO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004048566-A1
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                        cytotoxicity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUN-2004
                                                                                                                                               ADQ27128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NATO/) 1
(SAIG/) (TEIK/) 7
(NAIT/) 1
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                     Gaps
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0
Score 14.8; DB 1; Length 19;
Pred. No. 1.6e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                  Ribonucleotide tailing; sequencing; amplification; primer; terminal transferase; polymerase chain reaction; ss.
                                                                                                                                                                 Oligonucleotide 20-mer primer for ribonucleotide tailing.
                                       3286 CAGGAGAATTAGATTTCT 3303
                                                                                                                                                                                                                                                                        96DE-01001385
                                                                                                                                                                                                                                                                                            96DE-01001385
                                                                                                                                                                                                                                                                                                              (MUEL/) MUELLER M W.
                                                                                                                                                                                                                                                                         16-JAN-1996;
                                                                                                                                                                                                                                                                                            16-JAN-1996;
                                                                                                                                                                                                                                                      17-JUL-1997
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short inhibitory RNA; siRNA; adipocyte; cell membrane; electroporating Mouse Myolc targeted short inhibitory RNA (siRNA) target DNA SeqID102,

(first entry)

09-SEP-2004

ADP48862
ID ADP4
XX
AC ADP4
XX
XX
DT 09-S
XX

ADP48862;

RESULT 1564

요

Gaps

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Indels

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Gastrointestinal defensin peptide(s) - useful as antimicrobial and anti-inflammatory agents and for detecting gastrointestinal disorders.
                                                                                                                                                                                                     Sequence 20 BP; 8 A; 0 C; 3 G; 8 T; 0 U; 1 Other;
                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                       sites on the adapter or on the cDNA primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CHIL-) CHILDRENS HOSPITAL PHILADELPHIA.
                                                         Disclosure; Col 11; 12pp; German
                                                                                                                                                                                                                                                 2815 GTATATGGTATATACA 2832
                                                                                                                                                                                                                                                                GTATATGTTATATAAA 19
                                                                                                                                                                                                                                                                                                     AAQ53223 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                        93WO-US004740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-00888232
                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                   16; Conservative
                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jones DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1993-405719/50.
                     WPI; 1997-365083/34
                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                 defensin genes.
                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
17-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                           WO9324513-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                          09-DEC-1993
      Mueller MW
                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bevins CL,
                                                                                                                                                                                                                                                                                                                    AAQ53223;
                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                   Matches
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The sequences given in AAQ93370-441 represent oligomers which are capable of hybridising to an invariant proximal or distal sequence of a single mucleotide polymorphism (SNP) site in equine DNA. These oligomers act as primers, where template-dependent extension of the primer by a single nucleotide base can be used to identify the polymorphic allele. These sequences may be used to in the analysis of 18 SNP's from 5 different horses. The analysis of SNP's is useful in determining identity, ancestry or prediposition to genetic diseases. The method may also be used to determine the linkage between two genetic traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Invariant; proximal; distal; single nucleotide polymorphism; SNP; equine; human; primer; template-dependent extension; identification; polymorphic allele; horse; identity; genotype; ancestry; parentage; prediposition; genetic disease; linkage; ss.
                                         PCR amplification was conducted to determine tissue expression of defensin genes. For amplification of defensin 5 related sequences an upstream sense primer was chosen from one defensin-related open reading frame (HNP63S) and the downstream antisense primer was chosen from the other (HS1261a). These primers were chosen so the amplification product would include an intron when the template was genomic DNA, a possible contaminant in a pool of cDNA. Control amplifications from the cDNA templates used primer pairs (HTUBs and HTUBa) from the alpha-tubulin sequence. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid mols, for identifying single nucleotide polymorphism sites and related methods, useful for genotyping human or horse genome(s) for analysis of pre-disposition to genetic traits, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Equine clone 595-1 5' proximal antisense strand SNP detection oligo.
                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 8 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
Claim 50; Page 72; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 6; Page 49; 129pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              655 AATGGCAGCAAGGTGGGC 672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AATGGCAGCAAGGATGGC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
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94US-00216538.
                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ93404 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MOLE-) MOLECULAR TOOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Knapp MR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-193812/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Equus caballus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9512607-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-DEC-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ93404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ93404,
  8
                                                                                                                                                                                                                                                          This 20-mer oligonuclectide primer is used in a novel method of amplifying a nucleic acid molecule involving ribonuclectide tailing of single stranded DNA in the persence of terminal transferase. This method can be used for sequencing RNA, especially the 5'-end and involves a 3' tail of 2 to 4 ribonuclectides being attached onto the target sequence with high efficiency. The first step in cloning and sequencing the unknown 5'-end of a specific mRNA molecule involves the synthesis of single stranded cDNA using a primer specific for a known region in the 3' region of the mRNA, A tail of two rGTP's is attached to the mRNA using terminal transferase and a specific double stranded DNA adapter is attached via a complementary 3' overhang (i.e. CC) in the presence of T4 DNA ligase. PCR amplification using a 5'-bictinylated adapter specific primer and a cDNA-specific nested primer including a desired restriction site produces a unique PCR product suitable for direct solid-phase sequencing or dideoxy sequencing after a cloning step using restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gastrointestinal defensin peptide, GID; pharmaceutical; Paneth cell; antimicrobial; anti-inflammatory; diagnosis; probe; contact disinfectant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence of sense primer HNP63s used to determine tissue expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                 Nucleic acid amplification after 3' tailing - by reaction with ribonucleotide in presence of terminal transferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
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Sequence 20 BP; 11 A; 7 C; 1 G; 1 T; 0 U; 0 Other;

vivlemore401-10.rng

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(first entry)
                                                                                                                Equus caballus.
                                                                                                                                                                                                        03-NOV-1993;
23-MAR-1994;
                                                                                                                                      WO9512607-A1
                                                                                                                                                                                    02-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-DEC-1995
                                                                                                                                                             11-MAY-1995
                                                                                                                                                                                                                                                                 Goelet P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1570
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequences given in AAQ93370-441 represent oligomers which are capable of hybridising to an invariant proximal or distal sequence of a single mucleotide polymorphism (SNP) site in equine DN. These oligomers act as primers, where template-dependent extension of the primer by a single nucleotide base can be used to identify the polymorphic allele. These sequences may be used to in the analysis of 18 SNP's from 5 different horses. The analysis of SNP's is useful in determining identity, ancestry or prediposition to genetic diseases. The method may also be used to determine the linkage between two genetic traits
                                                                                                                                                                                                                   Invariant; proximal; distal; single nucleotide polymorphism; SNP; equine; human; primer; template-dependent extension; identification; polymorphic allele; horse; identity; genotype; ancestry; parentage; prediposition; genetic disease; linkage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid mols. for identifying single nucleotide polymorphism sites and related methods, useful for genotyping human or horse genome(8) for analysis of pre-disposition to genetic traits, etc.
                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                         ö
                                                                                                                                                                                              Equine clone 595-1 3' distal sense strand SNP detection oligo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03;
0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03; ved 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 1 A; 1 C; 7 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                              2352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 6; Page 49; 129pp; English
                                                                     N
                                                                                                                             BP.
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94US-00216538
           ilarity 88.9%;
Conservative
                                                              19 GIGIGIGIGIATITGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2335 GTGTGTGTGTGTGTGC
                                               2335 GTGTGTGTGTGTGTGTGC
                                                                                                                                                                                                                                                                                                                                                                                                         (MOLE-) MOLECULAR TOOL INC.
                                                                                                                            AAQ93403 standard; DNA; 20
                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 16; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                Knapp MR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-193812/25.
          Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                             Equus caballus
                                                                                                                                                                                                                                                                                                    WO9512607-A1
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                                                                                                                                                                         20-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                   23-MAR-1994;
                                                                                                                                                                                                                                                                                                                         11-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                Goelet P,
                                                                                                                                                   AAQ93403;
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   Query Match
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AAQ93405 standard; DNA; 20

RESULT 1569 AAQ93405, AAQ93405

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Invariant; proximal; distal; single nucleotide polymorphism; SNP; equine; human; primer; template-dependent extension; identification; polymorphic allele; horse; identity; genotype; ancestry; parentage; prediposition; genetic disease; linkage; ss.
                                             Invariant; proximal; distal; single nucleotide polymorphism; SNP; equine; human; primer; template-dependent extension; identification; polymorphic allele; horse; identity; genotype; ancestry; parentage; prediposition; genetic disease; linkage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid mols. for identifying single nucleotide polymorphism sites and related methods, useful for genotyping human or horse genome(s) for analysis of pre-disposition to genetic traits, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
Equine clone 595-1 3' distal antisense strand SNP detection oligo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Equine clone 595-1 5' proximal sense strand SNP detection oligo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 8 A; 1 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6; Page 49; 129pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2826 ATATACATATATATAT 2843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
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94US-00216538.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MOLE-) MOLECULAR TOOL INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Knapp MR;
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Gaps

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2; Indels

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Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor segment. The present sequence is an example of a compound SSR primer rethod represents a modified annulative of a compound SSR primer assay, which is partic. useful for genome fingerprinting, i.e. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer directed primer, comprising a sequence repeat (SSN), and an adaptor segment. The present sequence is an example of a compound SSR primer. The method repersents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adaptor directéd primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; compound; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detection; polymorphism; perfect compound simple sequence repeat;
                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Compound simple sequence repeat primer (AT) 2.5 (GT) 6.5.
                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 0 C; 7 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                              genetic trait marking and germplasm comparisons
                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (DUPO ) DU PONT DE NEMOURS & CO E I.
                     Example 2; Page 84; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 1c; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2316 TCTGTGTGTGTGTGTG 2333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 88.9 es 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vogel JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-277795/28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                           The sequences given in AAQ93370-441 represent oligomers which are capable of hybridising to an invariant proximal or distal sequence of a single nucleotide polymorphism (SNP) site in equine DNA. These oligomers act as primers, where template-dependent extension of the primer by a single nucleotide base can be used to identify the polymorphic allele. These sequences may be used to in the analysis of 18 SNP's from 5 different horses. The analysis of SNP's is useful in determining identity, ancestry or prediposition to genetic diseases. The method may also be used to determine the linkage between two genetic traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Modified amplified fragment length polymorphism assay - for detection of polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                           Nucleic acid mols. for identifying single nucleotide polymorphism sites and related methods, useful for genotyping human or horse genome(s) for analysis of pre-disposition to genetic traits, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ed primer; genome; genetic; fingerprinting;
yment length polymorphism assay; microsatellite region;
marking; germplasm comparisons; compound; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detection; polymorphism; perfect compound simple sequence repeat;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compound simple sequence repeat primer (AT) 3.5 (GT) 6.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 10 A; 1 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                       Claim 6; Page 49; 129pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT30406 standard; DNA; 20 BP
                                                                                     93US-00145145.
94US-00216538.
                                          94WO-US012632.
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                                                                                                                                                       (MOLE-) MOLECULAR TOOL INC.
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                                                                                                                                                                                                      Knapp MR;
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                                                                                                                                                                                                                                                 WPI; 1995-193812/25.
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                                          02-NOV-1994;
                                                                                       03-NOV-1993;
                                                                                                               23-MAR-1994;
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11-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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RESULT 1571

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Matches

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RNaseH; RNA cleavage; DNA cleavage; hybridisation; protein kinase C gene; gene expression modulation; ras; raf; therapy; AIDS; atherosclerosis; infection; cell growth; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligo:nucleotide with RNaseH activity, which specifically hybridises to DNA or RNA - comprises 1st and 2nd sub:sequence(s) having 2'-O-CH2-CH2-O-CH3 and 2'-deoxy sugar moieties, useful for therapy or diagnosis.
                                                                                                                                            Chimeric 2'-0-methyl oligo for c-raf inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 16; Page 41; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                  Monia B, Altmann K,
 20 CTGCCAGGGGAGGAG 3
                                                                AAX36462 standard; DNA; 20
                                                                                                                     06-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                         07-FEB-1997;
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                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                  Cook PD,
                                                                                          AAX36462;
                                       RESULT 1574
                                                     AAX36462
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT27481-T27507 are human c-raf kinase antisense oligonucleotides used for the inhibition of raf expression. The oligonucleotides (ONS) are targeted to either coding region, start or stop signal or 5' or 3' untranslated region (UTR) mNNA encoding human c-raf. The ONS may be phosphorothioate linked and may contain modifications at the 2' position of the sugar molety. ONs are pref. complementary to either 3' or 5' UTRs, phosphorothioate linked and contain 2'-O-alkyl sugar modifications. The ONS are used to inhibit expression of human raf in partic. in conditions associated with hyperproliferation e.g. cancer, restenosis, and psoriasis
                                                                  Gaps
                                                                                                                                                                                                                                                                    Human c-raf kinase 3' untranslated region antisense oligonucleotide
                                                                                                                                                                                                                                                                                             Antisense; anti-proliferative; tumour; cancer; raf; oncogene; phosphorothioate; 2' sugar modification; psoriasis; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligo:nucleotide(s) targetted to nucleic acids encoding human raf capable of inhibiting raf expression, used in treatment of hyper:proliferative disorders.
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                                     Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= b
/note= "contain 2'-O-methyl modifications"
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/note= "opt. phosphorothioate linked"
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             Sequence 20 BP; 3 A; 0 C; 7 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10; Page 18; 65pp; English.
                                                                                          2316 TCTGTGTGTGTGTGTG 2333
                                                                                                                     3 rarardrererereres 20
                                                                                                                                                                                       AAT27506 standard; DNA; 20 BP
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                                      Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative (
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                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                           misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                               misc_feature
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Martin P;

97WO-US002043. 96US-0011620P

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This sequence is an example of an oligonuclectide of the invention, and is an inhibitor of c-raf expression. The invention relates to conjugance (A), which specifically hybridises to RNA or DNA, comprises a linear sequence of nuclectide units linked by phosphodiseter or phosphorochicate linkages, comprising a first subsequence having 2'-O-CC CH2-CH2-O-CH3 sugar moieties and a second subsequence having 2'-O-CC CH2-CH2-O-CH3 sugar moieties and a second subsequence having 2'-O-CC CM2-CH2-O-CH3 sugar moieties and as RNAseM activity for cleaving a complementary strand, can be used to modulate the expression of ras, rafe and protein kinase C genes, useful in the therapy of AIDS, atherosclerosis, bacterial or other infections, or to control aberrant cell growth in humans, animals or plants. (A) can also be used diagnostically, particularly when labelled, to detect overexpression of chagnostically, particularly when including imaging of tissue sections, and as a research reagent. (A) has increased binding affinity for complementary strands (attributable to the 2'-O-CH2-CH3-O-CH3 sugar moiety, which overcomes the loss of affinity caused by altered intersugar collinks), and increased resistance to nuclease (from the modified links and
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               links), and increased resistance to :
the 2'-O-CH2-CH2-O-CH3 sugar moiety)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative
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844 CTGCCAGCCGAGGAG 861

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AMTS9716-28 are preferred oligonucleotides which are targeted to mRNA concoding human raf and are capable of inhibiting raf expression. Compositions containing the oligonucleotides entrapped in sterically stabilised liposomes are claimed. The compsns. can be used for inhibiting the expression of human raf. They can be used for the treatment of mammalian cancer, partic. human cancer e.g. lung, stomach, renal, breast, laryngeal, pancreatic, colorectal cancer and malignant melanoma. In particular the compsns. can inhibit abnormal raf expression and retain anti-hyperproliferative activity after prolonged circulation in the bloodstream. They facilitate the reduction of accumulation of ONs in nontarget organs and a reduction of acute and chronic side effects during prolonged treatment. ONls, 19 and 21 are chimeric oligonucleotides with uniform phosphorothicate backbones, and substituted with methoxy at the 2, position of the sugar moiety as indicated above. ON19 is targeted to
                                    1.12
//tag= b
//note="2' position of the sugar moiety is substituted by methoxy"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human c-raf and dextran sulphate mRNA targetting oligonucleotide ON19.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Inhibiting human raf expression, partic. for treating cancer - using oligonuclectide targetted to mRNA encoding human raf entrapped in sterically stabilised liposome(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cancer; anionic polysaccharide; human; lung cancer; stomach cancer; renal cancer; breast cancer; laryngeal cancer; pancreatic cancer; colorectal cancer; malignant melanoma; tumour; ss.
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
  a
"phosphorothioate backbone linkages"
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Best Local Similarity 88.9
Matches 16; Conservative
                     /note=
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misc_feature
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19-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a primer for the coding sequence for the protein of the invention. The protein of the invention is a recombinant protein (A), that comprises at least one region corresponding to an immunoglobulin (Ig) variable region which enables the protein to recognise and specifically bind to an antigen, preferably human Fas, and has substantially no more immunogenicity in a human patient than a human antibody. The proteins are useful for treating autoimmune diseases, especially rheumatism (rheumatoid arthritis). (A) is based on a murine monoclonal antibody. As the protein lacks the constant region, it has substantially no more immunogenicity in the human patient than a human
                                                                                                                                   Fas; antibody; human; immunoglobulin; variable region; rheumatism; autoimmune disease; rheumatoid arthritis; therapy; CDR; heavy chain; complementarity determining region; PCR primer; amplify; ss.
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                                                                                               Primer SHF-15 for H chain of Fas specific antibody coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-Fas recombinant antibodies - useful for treating auto-immune diseases, especially rheumatoid arthritis.
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Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                         (first entry)
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Les 16; Conservative
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modified_base
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                                                       15-APR-1998
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                AAV03702
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WPI; 1998-559441/48.
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28-JAN-1999
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                                                                                                                                                                                                                                                                                                                                                                                  Composition for cancer treatment - comprising anionic polysaccharide, and oligo:nucleotide targetted to mRNA encoding human c-raf and dextran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A pharmaceutical composition has been developed comprising an oligonucleotide, targeted to human raf encoding mRNA, and an anionic polysaccharide. The present sequence represents a specifically claimed oligonucleotide for use in the composition. The composition can be used to treat mammalian cancer, especially human lung, stomach, renal, breast, laryngeal, pancreatic or colorectal cancer, or malignant melanoma. The anionic polysaccharide increases tumour uptake of the oligonucleotide, particularly an oligonucleotide targeted to human raf encoding mRNA
              /note= "Phosphorothioate backbone; optionally being substituted at the 2'-position of the sugar moiety by a methoxy group at positions 1 to 12"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prostate, disease, PS112 gene, detection, diagnosis, cancer, treatment, antibody, primer, ss.
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Stroupe SD;
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Kratochvil JD, Roberts-Rapp L, Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PS112 gene RT-PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 16; Page 15; 21pp; English.
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                                                                                                                                                                                                                                                                                                      Nicklin PL, Steward A;
                                                                                                                                                                                                                                                              (CIBA ) CIBA GEIGY AG.
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AAV26668
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AC AAV26666
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                                                                                                                                                                AAV26668-V26669 are RT-PCR primers used in the isolation of the PS112 gene which is used in a novel method of detecting the presence of a target PS112 polynucleotide in a test sample. The method can also be used to detect mRNA of PS112 in a test sample. The method can be used for diagnosis of prostate cancer, as the presence of PS112 is an indicator of prostate cancer. Antibodies against the polypeptides may be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New Bacillus stearothermophilus DNA polymerase I clones - with reduced 3'-to-5' exonuclease activity useful for automated DNA sequencing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAV67205-10 are used to create 5' truncated Bacillus stearothermophilus DNA polymerase I enzymes. The truncated enzymes have DNA synthesis activity but are deficient in 3'-to-5' exonuclease activity. The new DNA polymerases are useful for nucleic acid synthesis by primer extension reactions, including manual, cycle (especially automated) DNA sequencing or labelling. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3' PCR primer P5 used to create a 5' truncated DNA polymerase I enzyme.
Detecting a target PS112 polynucleotide - used for diagnosing prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nerase I enzyme; truncated; primer extension; DNA sequencing; exonuclease activity deficient; nucleic acid synthesis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 0 A; 9 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                markers, or to treat prostate cancer
                                                                                                    Example 8; Page 88; 104pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            921 CTTCTTCCTGTTCATCCT 938
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Best Local Similarity 88.9
Matches 16, Conservative
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vivlemore401-10.rng

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New 2'-0-modified oligo-nucleotide(s) - comprising nucleotide(s) comprising a 2'-aminoalkoxy or 2'-imidazolylalkoxy substituent, used for hybridisation to RNA or DNA.
                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present oligonucleotide exemplifies the oligonucleotides of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nuclease resistant; ribofuranosyl moiety; 2'-aminoalkoxy; tumour; 2'-imidazolylalkoxy; modulation; activity; AIDS; atherosclerosis; phosphorothioate; ss.
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0
                                         0.4%; Score 14.8; DB 1; Length 20;
llarity 88.9%; Pred. No. 1.7e+03;
Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      c-raf antisense chimeric oligonucleotide of the invention.
BP; 2 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "phosphorothioated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 31; Col 50; 48pp; English.
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91WO-US005720.
92US-00835932.
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                                                                                                                                        3651 CTTGCTTGCCTGCAGGGC
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(first entry)
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cook PD, Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-166721/14.
                                                                 Local Similarity
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modified_base
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Sequence 20
                                                                                       16;
                                                                                                                                                                                                                                                                                                                                                                                                20-MAR-2003
16-APR-1999
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05-MAR-1992
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                                           Query Match
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AAX15068/C

ID AAX15068/C

XX AAX15068

DT 20-MAR-:
DT 20-MAR-:
DT 16-APR-:
XX Nucleas

XW D'-imid

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The present invention describes a method for measuring the resistance of p53 mutant cancer cells to the cytotoxic effects of chemotherapeutic agents by testing a sample comprising p53 mutant cells or an extract from p53 mutant cells for the abundance of cyclin protein D1. AAX90360 to AAX90373 represent reverse transcription PCR primers used to amplify the human p53 gene. The method can be used to predict the response of human cancer cells to anticancer therapy agents which can be used to select the most appropriate therapy for patients suffering from cancer. High cyclin D1 levels or high cyclin D1 expression together with p53 mutation is p1 levels or high cyclin D1 expression together with p53 mutation is strongly associated with resistance to cis-daminedichoroplatinum (CDDP) in human cancer cells. The test may be used to detect radiation may be a vibale alternative to CDDP or if other cycloxic agents would be more suitable, e.g. may suggest that Taxol should be considered as an alternative therapy as it may not be sensitive to a combination of p53 mutation and cyclin D1 protein overexpression
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molecules, or abnormal or inappropriate expression of normal RNA molecules in organisms or cells. (Updated on 20-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; p53; reverse transcription; PCR primer; resistance; mutant; cancer; cyclin D1 protein; chemotherapy; cytotoxic; ss.
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                                                                                                                                                                                           Score 14.8; DB 1; Length 20;
Pred. No..1.7e+03;
0; Mismatches 2; Indels
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18.9%; Pred. No. 1.7e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                 Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                   844 CTGCCAGCCGAGGAG 861
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1581
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Genetic proximity, gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;

Homo sapiens.

primer; 88 Synthetic WO9934016-A2

08-JUL-1999

PTK 15 gene specific primer.

(first entry)

11-OCT-1999

AAZ18120;

97IL-00122793. 98IL-00126627. 98WO-IL000625

28-DEC-1998;

29-DEC-1997; 16-OCT-1998;

AAZ18120 standard; DNA; 20 BP.

AAZ18120,

vivlemore401-10.rng

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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell the pattern cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a pecifical formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is characterising cells, a fetus. They can also be used for determining the an individual e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ17803-218342 represent primers that can be used in the RT-PCR reactions to determine set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor
                                                                                                                                                                                                                                                                   Genetic proximity; gene expression; cell characterisation; homeobox gene; gene; defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 0.4%; Score 14.8; DB 1; Length 20; Local Similarity 88.9%; Pred. No. 1.7e+03; nes 16; Conservative 0; Mismatches 2; Indels
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19 CCACTTGCCACCCTGCAC 2
                                                                                                                   AAZ18098 standard; DNA; 20 BP
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                                                                                                                                                                                                                                 PTK 4 gene specific primer.
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                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                         AAZ18098;
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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a discond cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for detecting a selected genetic defect in an individual, e.g. a fetus. They can also be used for determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired containing the pattern of gene expression in a selected gene family. Sequences Aal71003-Z13042 represent primers that can be used to the RT-PCR reactions to determine the pattern of gene expression. The method uses a contained the pattern of gene expression. The containing the pattern of gene expression. The containing the pattern of gene expression. The containing the contains the pattern of gene expression. The containing the pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of gene expression. The containing the pattern of pattern of pattern of genes with the containing the pattern of pattern of pattern of genes and pattern of patter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
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18.9%; Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               superfamily genes or cadherin superfamily genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 43; 102pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-419113/35.
P-PSDB; AAY14655.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENE-) GENENA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vider B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1584
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AAZ18104/c

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Gaps

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1801 GACGICTGGICCTTIGGG 1818

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Query Match

Matches

18 GACGTGTGGTCCTTCGGG 1

AAZ18104;

08-JUL-1999.

Vider B;

primer; ss. Synthetic.

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Human, raf, diagnosis, abnormal proliferative state, hyperproliferation, cancer, psoriasis, blood vessel restenosis, c-raf kinase, antisense, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nuclease resistant; modified; deoxyfuranosyl moiety; therapy; infection; AIDS; atherosclerosis; tumour; c-raf; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "contains phosphorothioate linkages; optional 2' -methyl modification on some base pairs"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotides targeted to human raf mRNA useful for treating and diagnosing abnormal proliferative states and inhibiting raf expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention provides antisense oligonucleotides targeted to mRNA encoding human raf and capable of inhibiting raf expression. The antisense oligonucleotides are useful for treating and diagnosing abnormal proliferative states and hyperproliferation (e.g. cancer, psoriasis, or blood vessel restenosis), and inhibiting raf expression. Sequences AAZI1511-537 and AAZI1555-573 represent antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                 Human c-raf kinase antisense oligo ISIS # 7854.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chimeric antisense oligo for c-raf gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotides for human c-raf kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 844 CTGCCAGCCGAGGAG 861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Col 11; 29pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 CTGCCAGGGGAGGAGGAG
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95WO-US007111.
                                                                                                                                                                                                                                                                                                                             96US-00756806.
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                    05-NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 88.9°
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-527018/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Boggs RT, Monia BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
modified_base
                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                             26-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                        31-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                              31-MAY-1995;
                                                                                                                                                                                                                                          US595229-A
                                                                                                                                                                                                                                                                                    14-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5859221-A.
                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX05466;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1586
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  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (c) determining in the first cell and the second cell; (d) determining in a selected gene family; and (c) calculating of proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for determining the crising cells capable of expressing an lomeobox related determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AA217803-218342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, protein phosphatase genes, because a set of portein receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                        Genetic proximity, gene expression, cell characterisation, homeobox gene, genetic defect, reverse transcriptase polymerase chain reaction, RT-PCR, kinase gene, protein phosphatase, P450, steroid receptor, cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 8 A; 7 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 1;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       superfamily genes or cadherin superfamily genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 42; 102pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1817
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                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-IL000625
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AAZ18104 standard; DNA; 20
                                                                                                                              PTK 7 gene specific primer.
                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-419113/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENE-) GENENA LTD.
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                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                 WO9934016-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-DEC-1997;
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                                                                                     11-OCT-1999
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Gaps

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AAZ11536;

RESULT 1585 AAZ11536/c ID AAZ11536 XX AC AAZ11536

Matches

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vivlemore401-10.rng

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The present invention describes a method for the diagnosis of a cancerous or pre-cancerous state from the co-elevation of cyclin-dependent kinase 1 (CDK1) and CDK4 levels. The method may be used for the clinical diagnosis of cancerous or pre-cancerous cells. In addition the combination of targets may be used to screen for drugs that may specifically act on cancer cells. The combination of CDK1, CDK4 elevation and p53 mutation in combination form a complex target that is likely to be specific for cancerous cells. AAX90388 to AAX90401 represent reverse transcription PCR primer for the human p543 gene, used in an example from the present
                                                                                                                                                                                                                                                                                         Diagnosis of cancerous or pre-cancerous cells by monitoring the levels of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, p53; reverse transcription; PCR primer; cancer, cytotoxic;
signal transduction factor; mutant; cell cycle; apoptosis; chemotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human p53 gene reverse transcription PCR primer exon 7 antisense.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                 cyclin-dependent kinases 1 and 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2695 CCACTTCCCACCTGCCC 2712
                                                                                                                                                                                                                                                                                                                                                           Example; Page 12; 26pp; English.
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                                                              98GB-00003447.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 88.9
Matches 16, Conservative
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                                                                                                                                                                                                 Seabra L;
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                                                                                                                                                     LIVERPOOL
                                                                                                                                                                                                                                             WPI; 1999-432548/37.
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                                                              18-FEB-1998;
                                                                                                           .8-FEB-1998;
                                                                                                                                                     (UYLI-) UNIV
                                                                                                                                                                                                 Warenius HM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-JUL-1998;
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                  25-AUG-1999,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nvention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a nuclease resistant compound that hybridises with RNA or DNA. The compound comprises covalently-bound nucleosides that individually include a ribose or deoxyribose sugar portion and a base portion, where the nucleosides to the nucleoside form a mixed base sequence that is complementary to a RNA base sequence or to a DNA base sequence that is complementary to a RNA base sequence or to a DNA base sequence; and where at least 1 of the nucleosides includes a modified deoxyfuranosyl moiety bearing a 2'-substituent selected from cyano, fluoromethyl, thioalkoxyl, alkylsulphinyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl, alkylsulphonyl alkeneoxy groups. The nuclease resistant oligonucleotides (ONs) can bind to organisms having a disease characterised by the undesired production of a protein. They can be used in therapeutic or prophylactic treatment in organisms such as bacteria, yeast, protozoa, algae, plant and higher animal forms including warm-blooded animals. The ONs can also be used for treating infections, AlbS, atherosclerosis or tumours. The products can be used for detection and diagnosis. The ONs provide enhanced binding to targets. Increased binding of 2'-sugar modified sequence-specific ONs provides superior potency and specificity compared to phosphorus-modified ONs. The present sequence represents a chimeric antisense oligo for craf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; p53; reverse transcription; PCR primer; cancer; diagnosis; mutant; cyclin-dependent kinase; CDK; ss.
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                                                                                                                                                                                                                                                                                                                                                                                Nuclease resistant oligonucleotide analogues - having nucleosides including modified deoxyfuranosyl moiety bearing 2'-substituent to increase binding affinity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human p53 gene reverse transcription PCR primer exon 7 antisense.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 31; Col 51; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       844 CTGCCAGCCGAGGAG 861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTGCCAGGGGGGGGGGG 3
                                                                                                                             90US-00566977.
91WO-US005720.
92US-00835932.
92US-00854634.
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                                                              95US-00468037
                                                                                                        90US-00463358
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                         Cook PD, Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                      WPI; 1999-120005/10.
                                                                                                                                13-AUG-1990;
12-AUG-1991;
05-MAR-1992;
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                                                            06-JUN-1995;
                                                                                                           11-JAN-1990;
                    12-JAN-1999
                                                                                                                                                                                                 01-JUL-1992
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WPI; 1999-422071/36.

GB2334578-A

Synthetic.

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ઠ g AAX90397;

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Gaps

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Indels

agent. Applications include determining identity (genetic

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sensitivity of cancer cells to anti-cancer agents by measuring the mutational status, expression and /or function of signal transduction factors. The method, by measuring the resistance of cells to anti-cancer agents, is useful for selecting the most appropriate therapy for patients suffering from cancer. AAX90374 to AAX90387 represent reverse transcription PCR primer for the human p543 gene, used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A method has been developed for the detection of one or more single polymorphisms in the same sample. The method comprises: (a) hybridising at least one distinguishable interrogation oligomorelectide primer (IP) to one or more target nucleic acids (TNA), each IP being specific for a unique region in TNA with the 3'-end of IP immediately adjacent to a specific and unique target nucleotide (nt); (b) extending IP with a template-dependent polymerase in presence of at least one non-extendible nt (or analogue); and (c) determining, for each IP, the identity of the non-extendible nt or analogue incorporated, i.e. the complement of the target nt, so identifying each target nt. The method is used to identify a trait (specifically a genetic disease or condition) in nucleic acid from an animal, plant, bacterium, fungus, yeast, virus, viroid or other
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Equine allele polymorphic site of clone 595-1 3'-distal sequence #1.
                             Determination of sensitivity of cancer cells to anti-cancer agents
                                                                                                                                                                        The present invention describes a method for the determination of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 14.8; DB 1; Length 20;
38.9%; Pred. No. 1.7e+03;
ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
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                                                                                                  Example 1; Page 18; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2695 CCACTTCCCACCTGCCC 2712
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polymorphisms in the same sample. The method comprises: (a) hybridising at least one distinguishable interrogation oligonucleotide primer (IP) to one or more target nucleic acids (TNA), each IP being specific for a unique region in TNA with the 3'-end of IP immediately adjacent to a specific and unique target nucleotide (nt); (b) extending IP with a template-dependent polymerase in presence of at least one non-extendible to an analogue); and (c) determining, for each IP, the identity of the non-extendible in to analogue incorporated, i.e. the complement of the non-extendible of a reaget int, so identifying each target nt. The method is used to identify a trait (specifically a genetic disease or condition) in nucleic acid from an animal, plant, bacterium, fungus, yeast, virus, viroid or other heritable agent. Applications include determining identity (genetic fingerprinting); ancestry; predisposition to genetic diseases or conditions (e.g. cancer, asthma, blindness); presence or absence of desirable trait; diagnosis of disease (e.g. haemophilia or sickle-cell anaemia) and to relate polymorphisms to particular traits. Axx03995 to AXX04067 represent the invariant 5'-proximal and 3'-distal sequences of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting single nucleotide polymorphisms - by hybridisation with interrogation primer and extending this with non-extendible nucleotide or analogue which is then identified.
          fingerprinting); ancestry; predisposition to genetic diseases or conditions (e.g. cancer, asthma, blindness); presence or absence of desirable trait; diagnosis of disease (e.g. haemophilia or sickle-cell anaemia) and to relate polymorphisms to particular traits. AAX03996 to AAX04067 represent the invariant 5'-proximal and 3'-distal sequences of polymorphic sites of corresponding equine alleles
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                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                    2; Indels
                                                                                                                                                                                           Score 14.8; DB 1; Length Pred. No. 1.7e+03;
                                                                                                                                                       Sequence 20 BP; 1 A; 1 C; 7 G; 11 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                            2335 GTGTGTGTGTGTGTGC 2352
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                                                                                                                                                                                               Query Match
Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting single nucleotide polymorphisms - by hybridisation with interrogation primer and extending this with non-extendible nucleotide or analogue which is then identified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          anaemia) and to relate polymorphisms to particular traits, AAXON996 to AAXO4067 represent the invariant 5'-proximal and 3'-distal sequences of polymorphic sites of corresponding equine alleles
                                                                                                                                                                                                                                                                                                               Equine allele polymorphic site of clone 595-1 5'-proximal sequence #2
                                                                                                                                                                                                                                                                                                                                         Polymorphic site; equine allele; detection; identification; identity; ancestry; genetic disease; cancer; asthma; blindness; haemophilia; sickle-cell anaemia; ss.
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                                                         Length 20;
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Pred. No. 1.7e+03;
                                                      Query Match 0.4%; Score 14.8; DB 1; Length 2 Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                            Sequence 20 BP; 8 A; 1 C; 1 G; 10 T; 0 U; 0 Other;
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polymorphic sites of corresponding equine alleles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Boyce-Jacino MT;
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                                                                                                                   2826 ATATACATATATATAT 2843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Goelet P,
                                                                                                                                               18 ATATCAATATATATAT 1
                                                                                                                                                                                                                         AAX04030 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                       AAX04030;
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Best Local Similarity

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An method has been developed for the detection of one or more single
polymorphisms in the same sample. The method comprises: (a) hybridising
at least one distinguishable interrogation oligonocleotide primer (IP) to
one or more target nucleic acids (TNA), each IP being specific for a
unique region in TNA with the 3'-end of IP lummediately adjacent to a
cunique region in TNA with the 3'-end of IP lummediately adjacent to a
specific and unique target nucleotide (nt); (b) extending IP with a
cepture of the capendent polymerase in presence of at least one non-extendible
cent (or analogue); and (c) determining, for each IP, the identity of the
non-extendible nt or analogue incorporated, i.e. the complement of the
carget nt, so identifying each target nt. The method is used to identify
ca trait (specifically a genetic disease or condition) in nucleic acid
from an animal, plant, bacterium, fungus, yeast, virus, viroid or other
centric fingerprinting); ancestry; predisposition to genetic diseases or
conditions (e.g. cancer, asthma, blindness); presence or absence of
desirable trait; diagnosis of disease (e.g. haemophilia or sickle-cell
anaemia) and to relate polymorphisms to particular traits. AAX03996 to
AAX04067 represent the invariant 5'-proximal and 3'-distal sequences of
polymorphic sites of corresponding equine alleles
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Equine allele polymorphic site of clone 595-1 5'-proximal sequence #1.
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Pred. No. 1.7e+03;
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2335 GTGTGTGTGTGTGTGC 2352
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                                                                   19 Grérérérérerarriée 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ancestry; genetic diseas:
sickle-cell anaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcintosh T, Head S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-081288/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Equus caballus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9859066-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-DEC-1998.
                                                                                                                                                                                                                                                                                                                         AAX04028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                               RESULT 1592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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The specification describes methods for detection of DNA and RNA basepair mismatch cleavage products, which indicates the presence and sequence of target DNA and RNA. Detection of the target is enhanced by amplification through recycling targets by maintaining an assay temperature between the melting point of the target/probe duplex and that of the target/probe duplex and that of the target/product complex, in the presence of an amplifier. The methods are used to detect and quantify specific DNA and RNA targets, e.g. in genetic diseases or infectious agents. They can also be used to detect mutations. The methods of the invention recycle the targets to dramatically increase the sensitivity of the mismatch repair assay, and allows for detection of a million or fewer target DNA on RNA molecules. PCR primers AAX26623-24
Detection; basepair mismatch cleavage product; mutation; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; c-raf; nuclease resistance;
RNAse H strand cleavage; phosphorothioate; oligonucleotide therapeutic;
AIDS; atherosclerosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide used to inhibit c-raf gene expression.
                                                                                                                                                                                                                                                                                                                                            Detection of DNA and RNA mismatch cleavage products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               were used to exemplify the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                   US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 12; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2695 CCACTTCCCACCTGCCC 2712
                                                                                                                                                                                     UNIV MARYLAND BALTIMORE.
                                                                                                                                                                                                                                                                                 Shih JW, Highsmith WE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19 CCACTTGCCACCCTGCAC 2
                                                                                                                           98WO-US018776,
                                                                                                                                                        97US-0058419P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                  HIGHSMITH W E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
                                                                                                                                                                                                                                                                                                               WPI; 1999-243734/20
                                                                                                                                                                                                     HSU I.
SHIH J W.
                                                                                                                           10-SEP-1998;
                                                                                                                                                      10-SEP-1997;
                                                             WO9913108-A1
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08-NOV-1999
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                                                                                            8-MAR-1999
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                              Synthetic.
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                                                                                                                                                                                                                  (SHIH/)
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AAZ10294/c
                                                                                                                                                                                     (UYMA-)
                                                                                                                                                                                                                                      HIGH/)
                                                                                                                                                                                                                                                                                 HBu I,
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                                                                                                                                                                     Vaccine, eye disease, conventional trachoma, nonendemic trachoma, paratrachoma, inclusion conjunctivitis, genital disease, perihepatitis, nongonococcal uretritis, epidymitis, cervicitis, salpingitis, PCR primer, bartholinitis, pneumopathy, venereal lymphogranulomatosis, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAX36734-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion epidymitis, genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingfitis, perihapatitis, barkholinitis, peneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for amplification of human breast tumour cell line DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                        primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 4 A; 2 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 1598; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1384 AACATCATCAACCTGCTG 1401
                                                                                                                                                                                                                                                                                                                                                                                         97FR-00015041.
97FR-00016034.
98US-0107077P.
                                             AAZ03336 standard; DNA; 20 BP
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AAX26624 standard; DNA; 20
                                                                                                         07-OCT-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 88.9
                                                                                                                                                                                                                                                 Synthetic.
Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-371125/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                            27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-1997;
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04-NOV-1998;
                                                                                                                                                                                                                                                                                                                             10-JUN-1999
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                                                                            AAZ03336;
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                1593
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Matches
           AAX26624
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ð g ö

Gaps

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Length 20; 2; Indels

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and other nucleic acid sequences from the genome of Chlamydia pneumoniae and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarccidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584-AAY35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides that detect and modulate the expression of Jun N-terminal kinase proteins - useful for treating hyperproliferative diseases and inhibiting tumor growth in animals, and for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to antisense oligonucleotides that detect and modulate the expression of Jun N-terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding a JNK1, JNK2 or JNK3 protein, and which modulate expression of these proteins. The oligonucleotides are useful for modulating JNK protein. The oligonucleotides are also useful for modulating JNK protein of a protein that has been phosphorylated by a JNK protein, and the expression of a cellular protein that protein that passen also form pharmaceutical composition events. The oligonucleotides also form pharmaceutical compositions for treating animals with a hyperproliferative disease, and for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNKl; JNK2; JNK2; Cell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chemically modified sense control probe ISIS No: 14318.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaarde WA;
                                                                                                                                                                                                                        Sequence 20 BP; 5 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                       Score 14.8; DB 1;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein phosphorylation by these proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 4; Page 92; 190pp; English.
                                                                                                                                                                                                                                                                                                                         743 TICICICCTIGCACAACG 760
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                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX29342 standard; DNA; 20
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                                                                                                                                                                                                                                                                       Local Similarity 88.9
nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1597
                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                     The present sequence represents a phosphorothioate antisense oligonuclectide used to inhibit c-raf gene expression. The oligonuclectide is a gapped 2' modified oligonuclectide, whereby one part has at least two consecutive 2'-F (2'-H) nuclectides and the second part has at least five consecutive nuclectides with 2'-H sugar moieties. The modified oligonuclectide has increased nuclease resistance, and increased binding affinity for substrates. The oligonuclectide elicits RNase H strand cleavage of specific RNAs, Oligonuclectides of the invention are useful for the diagnosis, detection and treatment of conditions susceptible to oligonuclectide therapeutics (e.g. AIDS and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                       Oligonuclectides eliciting RNAase H activity useful for diagnosis and treatment of diseases e.g AIDS or atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                             atherosclerosis). (Updated on 20-MAR-2003 to correct PR field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer used to amplify Chlamydia pneumoniae polynucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genome sequence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 1884; Disclosure; 1912pp; English
                                                                                                                                                                                                                         Example 14; Col 24; 34pp; English
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 91US-00814961.
92WO-US011339.
94US-00244993.
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98US-0107078P
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                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-357842/30.
                                                                                                                                       WPI; 1999-539598/45
                                                                                                      Cook PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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     24-DEC-1991;
23-DEC-1992;
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                                    21-JUN-1994;
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                                                                                                      Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
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BP.

tumor growth in an animal

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Human cystatin C gene PCR primer SEQ ID NO:4.
                                                AAA39285 standard; DNA; 20
                                                                                              13-SEP-2000 (first entry)
                                                                      AAA39285;
                                   AAA39285,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to antisense oligonucleotides that detect and modulate the expression of Jun N-terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding a JNK1, JNK2 or JNK3 protein, and which modulate expression of these proteins. The oligonucleotides are useful for modulating JNK protein expression and cell cycle progression in cultured cells or animal cells. The oligonucleotides are also useful for modulating the phosphorylation
                                                                                                                                                                                                                                                                          Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNKl; JNK2; JNK2; Gell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides that detect and modulate the expression o
Jun N-terminal kinase proteins - useful for treating hyperproliferative
diseases and inhibiting tumor growth in animals, and for modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of a protein that has been phosphorylated by a JNK protein, and the expression of a cellular protein that promotes one or more metastatic events. The oligonuclectides also form pharmaceutical compositions for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          treating animals with a hyperproliferative disease, and for inhibiting
                                                                      Gaps
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                                              Query Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaarde WA;
                        Sequence 20 BP; 2 A; 7 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein phosphorylation by these proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mckay R, Dean N, Monia BP, Nero PS,
                                                                                                                                                                                                                                                        JNK2-specific probe ISIS No: 12560.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; Page 87; 190pp; English.
                                                                                              1678 GACTICGGGCTGGCCCGG 1695
                                                                                                                     1 cacririscicriscicisci 18
                                                                                                                                                                                 AAX29331 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                  98WO-US016488.
                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-00910629
                                                                                                                                                                                                                               10-JUN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumor growth in an animal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-181060/15.
                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                   WO9909214-A1
                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1999.
                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                          AAX29331;
                                                                                                                                                                     SXS
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The present invention describes a method (M1) for diagnosing Alzheimer's disease (AD) comprising determining the level and/or activity of a transcription and/or translation product of a cystatin C gene (N1) or a polymorphic variant (N1a) of N1. Also described are: (1) a method (M2) for diagnosing AD comprising determining the presence or absence of a polymorphism in a cystatin gene; (2) a kit for carrying out M1 or M2; (3) a method (M3) for treating or preventing AD comprising administering an agent (I) which affects the level and/or activity of N1 or N1a, or the transcription and/or translation products of N1 or N1a; (4) the agent (I) in (3); (5) a method (M4) for identifying (I) comprising: (4) the agent (I) and a transcription product of N1 or N1a; (b) contacting the sample with (I); and (c) comparing the activity and/or level of the substances with (I); and (c) comparing the activity and/or level of the substances defere and after step (b). The methods are useful for prevention, changons and treatment of AD. The present sequence represents a PCR primer for the human cystatin C gene, which is used in an example from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing Alzheimer's disease comprising determining the level and/or activity of a transcription and/or translation product of a cystatin C gene or its polymorphic variant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
Keyhole limpet, diagnosis, Alzheimer's disease, cystatin C, neuroprotective, nootropic, gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 3 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 37; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2940 TGGAGGGAGGCCCCAGGG 2957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1600
AAA48651/C
ID AAA48651 standard, DNA, 20 BP.
XC AAA48651
                                                                                                                                                                                                                                                                                                                                                         98US-0105458P.
98US-0107434P.
99EP-00101377.
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                                                                                                                                                                                                                                                                                                99WO-EP008023
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Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 TGGTGGGAGGCCCCATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nitsch R, Growdon J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-350829/30.
                                                                                                                                                                  WO200025138-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NITS/) NITSCH
                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                22-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                             23.OCT-1998;
06-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                26-JAN-1999;
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Gaps

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0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; tive 0; Mismatches 2; Indels

1695

1678 GACTICGGGCIGGCCCGG 20 GACTITGGCCIGGCCCGG

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16; Conservative

Query Match Best Local Similarity Matches 16; Conserva'

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A novel method for modulating cell adhesion molecule expression involves antisense inhibition of a tumour necrosis factor (TNF) alpha signalling molecules. In the method TNF alpha signalling molecules Ha-ras, c-raf and c-dun N-terminal kinase (JNK) 2 were inhibited by antisense oligonucleotides. In addition an antisense oligonucleotide to the cell adhesion molecule E-selectin was also examined. The present sequence is the JNK2 antisense oligonucleotide. The antisense oligonucleotides used in the method contained modifications, namely phosphorothioate linkages and 2 methoxyethoxy bases. Some C residues also had a 5 methyl modification. Inhibitors of the TNF alpha signalling molecules have antibacterial, immunosuppressive, antipactic, antidiabetic, antithiranmatory activity. The antisense inhibitors may be useful for the tracement of sepsis, rheumatoid arthritis, inflammatory, immune disease, inflammatory bowel disease, allergic contact dermatitis, psoriasis,
                                                                               Antisense; E-selectin; TNF alpha; cell adhesion;
tumour necrosis factor alpha; phosphorothioate; methoxyethoxy; sepsis;
rheumatoid arthritis; inflammatory; immune disease;
inflammatory bowel disease; allergic contact dermatitis; psoriasis;
diabetes; Grave's disease; allergic contact concer; antibacterial;
immunosuppressive; antipsoriatic; antidiabetic; antithyroid; cytostatic;
dermatological; antiallergic; Ha-ras; c-raf; c-Jun N-terminal kinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modulating cell adhesion molecule expression for treating immune or inflammatory diseases involves treating cell with specific inhibitor of Tumor Necrosis Factor alpha signaling molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "All bases are 2'-methoxyethoxy, additionally bases are m5c"
                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Phosphorothioate internucleotide linkage"
                                               Antisense oligonucleotide ISIS no.15354 to human JNK2 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of sepais, rheumatoid arthritis, inflammatory, inflammatory bowel disease, allergic contact dermatitis, diabetes, Grave's disease, allograft rejection and cancer
                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 36; Page 46; 100pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
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                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                  modified base
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                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-DEC-1998;
                20-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUN-2000
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OTHER

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pCR primers AAA64094-95 were used to amplify a cDNA fragment of a human prostate cancer associated gene, which is designated PS112. PS112 sequences are useful for detecting, diagnosing, staging, monitoring, prognosticating, preventing, treating, or determining the predisposition of an individual to disease and conditions of the prostate, such as prostate cancer, tumours and metastases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel methods for diagnosing prostate cancer by contacting test sample with target specific polynucleotide and detecting prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gapa
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human biallelic marker upstream amplification primer SEQ ID NO:7832.
                                                                                                                                                                                                          PCR primer for prostate cancer associated gene PS112 cDNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Friedman PN, Gordon J, Hodges SC, Klass MR, Cohen M;
Roberts-Rapp L, Russell JC, Stroupe SD, Yu H, Kratochvil JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human genome, biallelic marker; high density disequilibrium map;
genomic map; haplotype; phenotype; polymorphic base; genotyping;
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Pred. No. 1.7e+03;
                                                                                                                                                                                                                                    Prostate cancer associated gene, PS112, prostate disease;
prostate cancer; tumour; metastasis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 9 C; 0 G; 11 T; 0 U; 0 Other;
7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; Col 71-72; 50pp; English.
                           1678 GACTICGGGCTGGCCCGG 1695
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ID AAZ73476 standard; DNA; 20 BP.
                                                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
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                                                       20 GACTTTGGCCTGGCCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     associated polynucleotides.
                                                                                                                           AAA64094 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                          (ABBO ) ABBOTT
                                                                                                                                                                                                                                                                                                                                                                    38-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                08-OCT-1996;
                                                                                                                                                                                                                                                                                   Homo sapiens
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 16;
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                                                                                                                                                       AAA64094;
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                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "All bases are 2'-methoxyethoxy, additionally C
bases are m5c"
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Length 20;

Score 14.8; DB 1; Pred. No. 1.7e+03;

0.4%;

Query Match Best Local Similarity

Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;

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24-SEP-1999;
                                                    Lu Z,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kawasaki AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6005087-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-JAN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-AUG-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ48164;
                                                    Sun X,
                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1604
                                                                                                                                                                                                                                                                                                                                                                                                      AAZ48164,
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                                                                                                                                                                                                                                                                         AAZ65654 to AAZ69578 represent human biallelic markers from the present nursention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and dispossic methods, as well as the characterisation of the pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ò
                                                                                                                                                                                                                       Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis B virus; HBV; Hepatitis A virus; HAV; probe; detection; mutation; high-density gene chip; ss.
haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; rive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis B virus related oligonucleotide probe #187.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                Chumakov I;
                                                                                                                                                                                                                                                       Claim 9; Page 1900; 2745pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2974 CAGAGGACCAGGGCTTTT 2991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA79924 standard; DNA; 20 BP
                                                                                                                            98US-0082614P.
98US-0109732P.
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                                                                                                        99WO-IB000822.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                 map of the human genome
                                                                                                                                                                                                   WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis B virus.
                                                                                                                                                           (GEST ) GENSET
                    diagnosis; ss
                                            Homo sapiens
                                                              WO9954500-A2
                                                                                                        21-APR-1999;
                                                                                                                            21-APR-1998;
                                                                                                                                        23-NOV-1998;
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                                                                                   28-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA79924;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polyribonucleotide solid phase synthesis; diagnosis; hybridisation; protein production modulation; 2'-deoxyfuranosyl moiety; anti-HIV; antiarteriosclerotic; nuclease resistant; atherosclerosis; AIDS; abnormal cell proliferation; tumour formation; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                             High-density gene chip making process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 examples from the present invention
                                                                                                                                                                                                                                                                                                                               Example 1; Fig 15; 19pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1877 AGGAGCTCTTCAAGCTGC 1894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 AGGAGCTCTTCTAACTGC 1
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91WO-US005720.
92US-00835932.
92US-00854634.
95US-00468037.
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99CN-00114460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 88.9°
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cook PD;
                                                             (UYDO-) UNIV DONGNAN.
                                                                                                                                                                                                WPI; 2000-443233/39
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WPI; 2000-400096/34.

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The present invention describes nuclease resistant oligonucleotides (I) comprising 2'-fluoro modified ribofuranosyl nucleotides. (I) comprise covalently bound nucleotides where the nucleotides in comparise by: (a) internucleotide linkages such that the base portion of the nucleotides forms a mixed base sequence; and (b) at least one of the nucleotides forms a modified ribofuranosyl group bearing a 2'-fluoro substituent; provided that at least two of the nucleotides are 2'-fluoro substituent; provided that at least two of the nucleotides are 2'-fluoro substituent; provided that at least two of the nucleotides are 2'-fluoro constituent; provided that at least two of the nucleotides are 2'-fluoro substituent; provided that to taste target RNA, and inhibit its expression. (I) are resistant to nuclease degradation and hybridise with appropriate strength and fidelity to its target RNA/DNA. (I) are also useful as research agents diagnostic agents and as oligonucleotide thereaged in conjunction with AZT to treat AIDS patients. (I) have been cat used to omodulate the expression of the treated arteries whose activate form has been implicated in abnormal cell proliferation and tumour formation. (I) are also used to modulate the expression of protein phosphorous modified oligonucleotide duplexes containing methylphosphonates, phosphoramidates and phosphate triesters. The present an oligonucleotide used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Forward PCR primer for detection of microsatellite marker TNFRSF1B intron 4 polymorphic region.
Nuclease resistant oligonucleotides useful as research agents, diagnostic agents, and in the treatment of atherosclerosis and AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumour necrosis factor; TNF; TNF-R2; TNFRSF1B; PCR primer; tumour necrosis factor receptor superfamily member 1B; human; cardiovascular disease; coronary artery disease; non-insulin dependent diabetes mellitus; neuropathy in NIDDM; essential hypertension; hyperlipidemia; diabetic neuropathy; vasoprotective; antihypertensive; lipid-lowering; chromosome 1p36.2; p1S2814; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                         Example 31; Col 51; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               844 CTGCCAGCCGAGGAGAG 861
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-NOV-1999;
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A novel method for determining a predisposition in a subject to a complex polygenic disease involves assaying chromosome 1 for a genetic marker indicative of a predisposition to the disease. This method may be used for determining predisposition to cardiovascular disease, coronary artery disease, non-insulin dependent diabetes mellitus, neuropathy in NIDDM, essential hypertension, hyperlipidemia and diabetic neuropathy. The method can be used for testing an individual with a family history or in the early stages of a complex polygenic disease to ascertain the chance of developing hypertension, neuropathy or lipid disturbances such as high total cholesterol, high low density lipoprotein cholesterol, abnormal apolygenic disease to ascertain the chance coplygenic disease disposition has been identified the subject can be treated to prevent or reduce the disease or delay its onset. The genetic marker used in the method is DIS2834 and includes a CA repeat region in intron 4 of the tumour necrosis factor receptor superfamily member 1B (TNPRSTB) gene. The marker is located at chromosome 1936.2. The present sequence is the forward PCR primer used for detection of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to antirheumatic agents which comprise as
                                    Method for diagnosing a predisposition to a complex polygenic disease e.g. coronary heart disease, hyperlipidemia and non-insulin-dependent diabetes mellitus comprises assaying chromosome 1 for a genetic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         active ingredients an immunoglobulin M (IgM) procein. The IgM protein does not include a J segment, has apoptosis inducing activity, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-human Fas humanizing antibody-containing antirheumatic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-human Fas antibody CH11 H chain cDNA specific primer SHF-15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  microsatellite marker TNFRSF1B intron 4 polymorphic region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2316 TCTGTGTGTGTGTGTG 2333
                                                                                                                     Claim 30; Page 4; 50pp; English
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consists of a light and heavy chain polypeptide produced synthetically. The agents of the invention exhibit antirheumatic and immunosuppressive activity and can be used to treat autoimmune diseases, especially rheumatism. The IgM molecule used in the invention has human Fas-antigen binding properties. Included in the invention are nucleotide sequences of the invention are nucleotide sequences of corresponding protein sequences (see AAB12913-1812918 and AAB12919), and corresponding protein sequences (see AAB12913-1812918 and AAB12919), and nucleotide sequences of the humanised anti-human Fas Ig CH11 (see AAB12902-AAB206) and protein sequences (see AAB12908-B12910). Also included are anti-human Fas antibody CDP peptides (AAB12902-B12907). Primers specific for the anti-human Fas antibody, light, heavy and kappa chains used in the invention are represented by sequences AAA78213-A7826. Primers used for sequences AAA78213-A78318 and AAA78335-A78313, while humanised anti-Fas Ig DNA sequencing primers are sequences AAA78212-A78313 and AAA78331-A78312-A78331 and AAA78331-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A78313-A
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Matches 16; Conservative
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GTGTACAGTGACGCACAG 621 604 ò 셤

Human fra-1 mRNA antisense oligonucleotide ISIS 109078. AAC60587 standard; DNA; 20 (first entry) 31-JAN-2001 AAC60587; RESULT 1607 AACGOS87/C
AACGOS87/C
AACGOS87/C
AACGO
AAC

BP

Human, fra-1, antisense oligonucleotide, phosphorothioate, cytostatic, antiinflammatory, 2'-methoxyethyl wing, 2'-MOB wing, infection, cancer,

Homo sapiens Synthetic.

US6124133-A

26-SEP-2000

15-0CT-1999;

99US-00418641. 15-OCT-1999;

99US-00418641.

(ISIS-) ISIS PHARM INC

Taylor JK, Cowsert LM;

WPI; 2000-601552/57.

Novel antisense compound 8-30 nucleobases in length targeted to human fra -1 and which specifically hybridizes with and inhibits the expression of human fra-1, useful for modulating the expression of fra-1 in cells. The present sequence is one of a large number of antisense oligonucleotides which are targeted to nucleic acids encoding fra-1. The sequences may be oligodeoxyribonucleotides or chimeric oligonucleotides containing a central gap region consisting of ten 2'-deoxynucleotides, which is flanked on both sides by 2'-methoxyethyl (2'-MOE) wings. The oligonucleotides have a phosphorothioate backbone and the cytidine Claim 3; Col 42; 38pp; English.

0.4%; Score 14.8; DB 1; Length 20;

Query Match

Sequence 20 BP; 2 A; 7 C; 7 G; 4 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel methods for reducing apoptosis comprising contacting cells with antisense oligonucleotides, useful for treating apoptotic disorders, e.g.
residues in the 2'-MOE wings are 5-methylcytidines. The fra-1 antisense oligonucleotides are useful for inhibiting the expression of fra-1 in human cells or tissues. They can be used for diagnostics, therapeutics, prophylaxis and as research reagents and in kits. Use of the antisense compounds may also be useful prophylactically, e.g. to prevent or delay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense; gene therapy; JNK2 protein; apoptosis; cancer; cellular hyperproliferation; Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney;
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                                                                                                                                                                Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
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                                                                                            infection, inflammation or tumour formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JNK antisense oligonucleotide ISIS #14318.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diabetes; Jun N-terminal kinase; ss.
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                                                                                                                                                                                                                                            1123 ACGCTGGCCAATGTCTCC 1140
                                                                                                                                                                                                                                                                               20 ACCCTAGCCAATGTCTCC 3
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                                                                                                                                                                  Query Match
Best Local Similarity
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Human, c-raf, protein kinase, antisense oligonucleotide, cancer, signal transduction; hyperplasia, pulmonary fibrosis; angiogenesis; psortiasis, atherosclerosis; macch muscle cell proliferation; stenosis; restenosis; inflammatory disorder; tissue graft rejection; endotoxin shock; glomerular nephritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotides targeted to nucleic acid molecule encodi
human raf useful for diagnosis, treatment of raf-associated cell
proliferative conditions such as cancer, psoriasis or blood vessel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             methoxyethoxy modification. Also, optionally phosphodiester or phosphothioate backbone"
                                                                                                                                            c-raf kinase antisense oligonucleotide #35 (Isis #7854).
AAA73514 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-00250856.
95WO-US007111.
96US-00756806.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-00143214.
                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boggs RT, Monia BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-531424/48.
                                                                                                                                                                                                                                                                                                                                                                                                             modified base
                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-NOV-1996;
                                                                                             28-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US6090626-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        restenosis.
  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to antisense oligonucleotides (AAC62844-C63000, AAA6099-A96099 and AAA07993) that hybridise specifically to a nucleotide encoding a Jun N-terminal kinase (JNK2) protein, resulting in decrease of JNK2 expression and leading to induction of apoptosis. The present sequence is one such antisense oligonucleotide. The oligonucleotides of the present invention are useful for treating diseases or conditions with reduced apoptosis, e.g. cancer and cellular hyperproliferation. The oligonucleotides may also be used to increase the stimulation of apoptotic proteins, e.g. for treating Alzheimer's or Parkinson's disease, amylotrophic lateral sclerosis, retinitis, pigmentosa, epilepsy, mycoardial infarction, stroke, obstructive jaundice, polycystic kidney and diabetes. The present sequence may have a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel methods for reducing apoptosis comprising contacting cells with antisense oligonucleotides, useful for treating apoptotic disorders, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense; gene therapy; JNK2 protein; apoptosis; cancer; cellular hyperproliferation; Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney; diabetes; Jun N-terminal kinase; ss.
                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
  Pred. No. 1.7e+03;
); Mismatches 2; indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    JNK antisense oligonucleotide ISIS #12560.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nero PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 133; 160pp; English.
                                                                      1695
                                                                                                    1 GACTTTGGCCTGGCCCGG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mckay R, Dean NM, Monia BP,
                                                                                                                                                                                                                                            踞
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-APR-2000; 2000WO-US008880.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00287796
88.9%;
                                                                      1678 GACTICGGGCTGGCCCGG
                                                                                                                                                                                                                                            AAC62874 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      phosphorothicate backbone
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-638427/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200059549-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-APR-1999;
                                                                                                                                                                                                                                                                                                                                         06-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-OCT-2000
                                                                                                                                                                                                                                                                                         AAC62874;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer
                                                                                                                                                                                          AAC62874/C
AAC68/XX
AAC68
XX
AAC66
AAC6
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/mod_base= OTHER /note= "All or some nucleotides are optionally with 2'-

base= OTHER

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1. .20 /*tag=

Location/Qualifiers

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c-raf is a serine-threonine-specific protein kinase and is thought to play a fundamental role in signal transduction, and cell proliferation control. The present sequence is an antisense oligonucleotide. This sequence is targeted to human c-raf gene, resulting in c-raf expression inhibition. The present sequence may be useful for treating and raf-ssociated cell hyperpoliferation conditions such as cancer, hyperplasias, pulmonary fibrosis, angiogenesis, psoriasis, atherosclerosis and smooth muscle cell proliferation in blood vessels egg stenosis or restenosis following angioplasty. Also, the present sequence may be useful for treating inflammatory disorders such as tissue graft rejection, endotoxin shock and glomerular nephritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
Claim 31; Col 10; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           844 CTGCCAGCCGAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          creccaeegageage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS06848 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS06848
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Gaps

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1678 GACTTCGGGCTGGCCCGG 1695

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Local Similarity 88.9

Matches

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GACTTTGGCCTGGCCCGG

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RESULT 1610 AAA73514/c Human bcl-6 phosphorothioate antisense oligonucleotide, SEQ ID NO:76.

(first entry)

23-FEB-2001

transcriptional repressor; germinal centre formation;

Human; bcl-6; Th-2 mediated

Homo sapiens

31-OCT-2000,

US6140125-A.

Th-2 mediated antibody affinity maturation; apoptosis regulator; chromosome 3027; Immboma; acute lymphoblastic leukaemia; post-transplant lymphoproliferative disorder; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.

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AASO6832-AASO6897 represent part of a polynucleotide sequence encoding for novel human protein kinases where a single nucleotide polymorphism (SNP) has been identified. The SNP occurs at the last position of the present sequence. The sequences are described relating to the invention of the protein kinase for the sequences are described relating to the invention of protein kinase for a members of the tyrosine or concing protein kinase (PTK and STK) families. The polynucleotides expression. STK families. The polynucleotides on concing protein kinase and the polypeptides may be used in the inappropriate kinase expression. For example, they may be used to treat cancers (especially cancers of haematopoidetic origin), cardiovascular cancers (especially cancers of haematopoid arthritis), neurological disease (e.g. atherosclerosis), metabolic disorders (e.g. diabetes), immune related diseases (e.g. rheumatoid arthritis), neurological disease (e.g. schizophrenia), neurodegenerative disorders (e.g. diabetes), for parkinson's disease (e.g. HIV) and reproductive disorders (e.g. infertility). Additionally, polynucleotides encoding protein kinases may be used for gainst the protein kinases and in assays to identify modulators of
                                                                                                                                   Human; protein kinase; PTK; STK; cancer; cardiovascular disease; SNP; metabolic disorder; immune related disease; neurological disorder; neurodegenerative disorder; inflammatory disorder; infectious disease; reproductive disorder; gene therapy; single nucleotide polymorphism; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding human kinase polypeptides, useful for preventing diagnosing and/or treating e.g. cancer, immune, cardiovascular and neuronal-associated diseases, and microbial infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sudarsanam S, Martinez R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 2 A; 6 C; 7 G; 4 T; 0 U; 1 Other;
                                                                                               SNP containing protein kinase DNA sequence #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein kinase expression and activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8B; Page 330; 433pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Manning G,
                                                                                                                                                                                                                                                                                                                                                              22-NOV-2000; 2000WO-US032085
                                                                                                                                                                                                                                                                                                                                                                                                      99US-0167482P
                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Whyte D,
Clary D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-343950/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                           (SUGE-) SUGEN INC.
                                                                                                                                                                                                                                                                               WO200138503-A2
                                                                                                                                                                                                                                                                                                                                                                                                    24-NOV-1999;
                                                                                                                                                                                                                                           Homo sapiens
                                                          12-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plowman GD,
                                                                                                                                                                                                                                                                                                                    31-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Flanagan P,
                    AAS06848;
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Antisense compounds which specifically hybridize with and inhibit human bcl-6 expression, useful for treating bcl-6 related disorders, and preventing or delaying inflammation or tumor formation.

99US-00418640.

15-OCT-1999; 15-OCT-1999;

Cowsert LM;

Taylor JK,

WPI; 2001-048959/06.

(ISIS-) ISIS PHARM INC

Example 15; Col 43-44; 42pp; English.

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Sequences AAC81144-C81223 represent antisense oligonucleotides targetted to the human bcl 6 gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human bcl-6 mRNA, and were analysed for their effect on bcl-6 mRNA levels by quantitative real-time PCR. Bcl-6 (also known as B-cell CLL/ lymphoma 6, and finger protein 51 and LAZ3) is a sequence-specific DNA-binding transcriptional repressor. The bcl-6 gene is expressed in germinal centre B- and T- cells and is required for germinal centre formation and Th-2 mediated antibody affinity maturation. Bcl-6 may also play a role in the region which undergoes a high frequency of translocation events. Such crossomal translocations can result in abbrarant forms of Bcl-6, which are strongly implicated in the pathogenesis of several types of lymphoma, and have also been reported in acute lymphoblastic leukaemia and post-transplant lymphoplastic disponders. The algonomicleotides of the conditions associated with aberrant forms of bcl-6, such as lymphomas, also expended with aberrant forms of bcl-6, such as lymphomas, also expended by a post-transplant lymphoblastic leukaemia and post-transplant lymphoblastic leukaemia and post-transplant lymphoblastic leukaemia and post-transplant lymphoplastic leukaemia and post-transplant lymphoplastic leukaemia and post-transplant lymphoplastic leukaemia and post-disposis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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0
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 0 C; 2 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3458 AAGTITATATATATCTAT 3475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 AGGITTATATATATTAT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF72931 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%;
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF72931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1613
BXXXXX
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Gaps ö

Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03; 0; Mismatches 2; Indels

0; Mismatches

0.4%; 88.9%;

Query Match Best Local Similarity 88.99 Marches 16; Conservative

520 18

503 TGGACGTGCTGGAGCGCT rédecériecresadeceer AAC81210 standard; DNA; 20 BP.

RESULT 1612 AAC81210 ID AAC81210 XX AC AAC81210

AAC81210,

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Homo sapiens.

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nucleobases in length, where (I) inhibits expression of daxx (also known as Fas binding protein, CENP-C binding protein, dap6 for death associated protein, and EAP for Ets-1 associated protein). (I) has cytostatic and antiinflammatory activity, and can be used in antisense therapy and as a modulator of daxx. (I) is useful for inhibiting the expression of daxx in therapeutics for the treatment of diseases associated with the expression of daxx, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation and as research reagent. The present sequence represents an inhibitory human daxx antisense phosphorothioate oligonucleotide which
                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel antisense compounds capable of modulating expression of daxx useful for diagnosis, prophylaxis and treatment of diseases associated with expression of daxx.
                  Antisense oligonucleotide; daxx; inhibition; phosphorothioate;
Fas binding protein; CENP-C binding protein; dap6; EAP; cytostatic;
antiinflammatory; death associated protein 6; Ets-1 associated protein;
infection; inflammation; tumour formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention describes an antisense compound (I) up to 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Col 42; 59pp; English
                                                                                                                                                                                                                                                                           24-JAN-2000; 2000US-00490692.
                                                                                                                                                                                                                                    24-JAN-2000; 2000US-00490692
                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                         Dean NM, Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-217744/22.
                                                                                                                                                          US6180353-B1.
                                                                                                                   Homo sapiens
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                                                                       y Match 0.4%; Score 14.8; DB 1; Length 20; Local Similarity 88.9%; Pred. No. 1.7e+03; nes 16; Conservative 0; Mismatches 2; Indels
used in the exemplification of the present invention
                                                                                                                                                       1169 GGCAGCTGTCTCGGGCCC 1186
                                                                                                                                                                                          18 GGTCCTGTCTCGGGCCC 1
                                                                         Query Match
                                                                                                                Matches
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AAF59860 standard; DNA; 20 BP

(first entry)

04-MAY-2001

Human

RESULT 16
AAF59860/
ID AAF59860/
XX
AC AAF5;
XX
DT 04-M
XX
KW Human
XX
KW isozy
KW Calc;
KW Jun b
KW T-ce
KW Jun b
KW GYCOI
KW GY

AAF59860

Sequences AAF59817-AAF59896 represent phosphorothioate 2'-WOE gapmer antisense targetted to the human protein kinase C-theta gene, which inhibit its expression. The antisense oligomucleotides were designed to target different regions of the human protein kinase C-theta RNA, and were analysed for their effect on protein kinase C-theta RNA, and were analysed for their effect on protein kinase C-theta RNA, levels by quantitative real-time PCR. Protein kinase C-theta (also known as PKC-theta, PKCT, PKKCT, nPKC-theta and PKKCO) is one of several protein kinase C isozymes and is ubiquitously expressed, with the highest levels kinase C isozymes and is ubiquitously expressed, with the highest levels can a calcium-independent fashion, and it is involved in a variety of in a calcium-independent fashion, and it is involved in avariety of the JNK/SAPK (Jun N-terminal Kinase/Atress-activated protein kinase) pathway. Protein kinase C-theta is also involved in T-cell signalling pathways, cell cycle control, cellular activation, API transcription pathways cell cycle control, cellular activation, and angiogenesis on wound repair. It is additionally involved in insulin signalling and is thought to play a role in the development of diabetes in humans. The coligonucleotides of the invention are useful for diabetes in humans. The and treatment of conditions associated with protein kinase C-theta

expression, such as inflammation, cancer, wound healing disorders and

Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;

0.4%;

Ouery Match Best Local Similarity 88.9 Matches 16; Conservative

Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human protein kinase C-theta useful for inhibiting expression of human protein kinase C-theta in human cells.

99US-00429322 99US-00429322

26-OCT-1999; 20-FEB-2001.

(ISIS-) ISIS PHARM INC

WPI; 2001-210378/21.

Bennett CF,

Claim 3; Col 43-44; 40pp; English

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Gaps

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Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03; 0; Mismatches 2; Indels

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Fibroblast growth factor 23; FGF-23; injury; placental cell; ulcer; congenital defect; fertility; abnormal growth; thymus function; leukemia; lymphoma; autoimmune disease; proliferative disorder; differentiation disorder; central nervous system disorder; infarction; Parkinson's disease; Alfheimer's disease; inflammation; intestinal wound; motility disorder; absorption disorder; stroke; congenital malformation; ischemic vascular disease; myocardial ischemia;
                                                                                                                                                                                                                                                                                                                                                                                         PCR primer for human fibroblast growth factor 23 (FGF-23) cDNA.
1872 TGTGGAGGAGCTCTTCAA 1889
                                                                                                                                                                                                            BP.
                                                                                                                                                                                                            AAH75019 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                   29-OCT-2001
                                                                                                                                                                                                                                                                        AAH75019;
                                                                                                                                                RESULT 1615
                                                                                                                                                                                   AAH75019/
                                                                                                                                                                                                            δ
                                                                                                                                                                                                                                                                                                                                protein kinase C-theta; PKC-theta; PKCT; PRKCT; nPKC-theta; PRKCQ;
                                                                                                                                                                                                                                                                    Human protein kinase C-theta antisense oligonucleotide, SEQ ID NO:53
                                                                                                                                                                                                                                                                                                                                                isozyme; serine/threonine protein kinase; signal transduction; calcium-independent function; JNK/SAPK pathway upstream activator; Jun N-terminal kinase/etress-activated protein kinase;
Jun N-terminal kinase/etress-activated protein kinase;
Jun Signalling pathway; cell cycle control; cellular activation;
API transcription factor activation; AIDS aetiology; apoptosis; cytoskeletal arrangement; proliferation; wound healing disorder; angiogenesis; insulin signalling; chromosome 10p15; phosphorothioate; 2'-MOB gapmer; ss. ancer; inflammation; diabetes;
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CC Growth factor 23 (FGF-23). Human FGF-23 is useful for treating a patient growth factor 23 (FGF-23). Human FGF-23 is useful for treating a patient suffering from traumatic injury or a condition characterized by dysfunction of or injury to skin cells, a condition characterized by inadequate function of placental cells (e.g. congenital defects, fertility, or abnormal growth), a condition characterized by inadequate function of the thymus (e.g. lewkemia, lymphoma, autoimmune disease, function of the thymus (e.g. lewkemia, lymphoma, autoimmune disease, proliferative disorder of the thymus, or differential nervous system disorder of the thymus, or a condition characterized by central nervous system disorder (e.g. Parkinson's disease or Alzehimer's disease). The human FGF-23 polynucleotide and polypeptide are also useful in the treatment of congenital malformations of the intestinal wounds, ulcers, inflammation, injuries and surgical anastromoses, motility and absorption disorders, or resulting ischemic and stand diseases (e.g. mycocardial ischemial infarction, peripheral vascular disease, renal artery disease, stroke) and musculoskeletal disease characterized by loss of function, inadequate colls, hone cells, hone cells, bone cells, carporting cells (e.g. mycolle or arkhing merching or death of skeletal muscle cells, hone cells, bone cells, carporting cells (e.g. mycordial disease)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated nucleic acids encoding the human and murine fibroblast growth factor 23, useful in the treatment of a condition characterized by inadequate function of placental cells (e.g. congenital defects) and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human nucleolin phosphorothioate antisense oligonucleotide, SEQ ID NO:11.
peripheral vascular disease; renal ártery disease; bone disease;
musculoskeletal disease; skeletal myopathy; arthritis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human nucleolin, P92; C23; phosphoprotein, ribosome biogenesis,
ribosome transport, cytokinesis, nucleogenesis, cell proliferation,
cell growth, transcriptional repression; replication;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cells (e.g. skeletal myopathies, bone disease, or arthritis)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 14.8; DB 1; Length 20; 18.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 7 A; 8 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 41; 77pp; English.
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                                                                                                                                                                                               07-MAR-2001; 2001WO-US007469.
                                                                                                                                                                                                                                        08-MAR-2000; 2000US-0187854P.
18-SEP-2000; 2000US-023368P.
05-DEC-2000; 2000US-0251649P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  825 CTCTGCGTGGCTGGTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 crcrcacrccrccrccr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC92561 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               thymus (e.g. leukemia).
                                                                                                                                                                                                                                                                                                                                                                                               Itoh N, Kavanaugh MW;
                                                                                                                                                                                                                                                                                                                             (CHIR ) CHIRON CORP. (KYOU ) UNIV KYOTO.
                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-522948/57.
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                                                                                                            WO200166596-A2.
                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAR-2001
                                                                                                                                                       13-SEP-2001
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ID AAC
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Sequences AAC92560-C92639 represent antieense oligonucleotides targetted to the human nucleolin gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human nucleolin meNA, and were analysed for their effect on nucleolin meNA.

Coligonucleotides were designed for their effect on nucleolin meNA levels by quantitative real-time PCF. Nucleolin (also known as P92 or C21) is the most abundant nucleolar phosphoprotein in actively growing cells. Nucleolin primarily participates in ribosome blogenesis and transport of ribosomal components, being able to transiently bind to previous some in the nucleolar being able to transiently bind to previous some in the nucleolar via a ribonucleoprotein consensus sequence. Thewever, it has also been shown to be involved in cytokinesis, coll proliferation and growth, transcriptional repression, replication, signal transduction, and chromatin decondensation. Nucleolin is a member of the Ag-NOR (active ribosomal gene located in the nucleolar corganiser region) family of proteins which are markers of active ribosomal genes, and whose expression is associated with the prediction of tumour growth rate. The presence of antibodies against nucleolin are associated with systemic connective tissue diseases such as systemic conception are useful for diagnosis, conversion and treatment of conditions associated with nucleolin are prevention and treatment of conditions associated with nucleolin are prevention and treatment of conditions associated with nucleolin and prevention and treatment of conditions associated with nucleolin and convention and treatment of conditions associated with nucleolin and appropriated with an analysis and solectodermaliated with nucleolin and the presence of the invention are useful for diagnosis, and active associated with nucleolin and and active the presence of antibodicated with nucleolin and active the presence of a nucleolin and appropriation and appropriated with an active formation and presence asso
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense compound targeted to human nucleolin which specifically hybridizes with and inhibits the expression of human nucleolin, useful for modulating the expression of nucleolin in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression, such as tumour formation, immune disorders and inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
signal transduction; chromatin decondensation; Ag-NOR family; nucleolin antibody; systemic connective tissue disease; SLE; systemic lupus erythematosus; scleroderma-like chronic graft versus host disease; expression inhibition; tumour formation; cancer; inflammation; immune disorder; phosphorothioate; antisense oligonucleotide; ss.
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; Col 41-42; 41pp; English.
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AAC92593 standard; DNA; 20 BP.
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Local Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51 GCGGCTGCAGGTGCTGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cowsert LM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                     Homo sapiens
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modified_base
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                                                         Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequences AAC92560-C92639 represent antisense oligonucleotides targetted to the human nucleolin gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human nucleolin mRNA, and were analysed for their effect on nucleolin mRNA is not were analysed for their effect on nucleolin mRNA levels by quantitative real-time PCR. Nucleolin (also known as P92 or C23) is the most abundant nucleolar phosphoprotein in actively growing cells. Nucleolin primarily participates in ribosome biogenesis and cells. Nucleolin primarily participates in ribosome biogenesis and ribosome in the nucleolar will a ribonucleoprotein consensus sequence. However, it has also been shown to be involved in cytokinesis, nucleogenesis, cell proliferation and growth, transcriptional repression, replication; signal transduction, and chromatin decondensation. Nucleolar or sa member of the Ag-NOR (active ribosomal gene located in the nucleolar corganiser region) family of proteins which are markers of active ribosomal genes, and whose expression is associated with hystemic connective tissue diseases such as systemic lupus erythemactosus (SIE) and selecotated with nucleolin are associated with systemic connective tissue diseases such as systemic disease. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with nucleolin
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                                                                                                                                                                                                                                                                                                                                                        Novel antisense compound targeted to human nucleolin which specifically hybridizes with and inhibits the expression of human nucleolin, useful for modulating the expression of nucleolin in cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression, such as tumour formation, immune disorders and inflammation
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 ribosome transport; cytokinesis; nucleogenesis; cell proliferation;
                                                     systemic lupus erythematosus;
scleroderma-like chronic graft versus host disease;
expression inhibition; tumour formation; cancer; inflammation;
immune disorder; phosphorothioate; antisense oligonucleotide; ss.
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           cell growth, transcriptional repression, replication, signal transduction, chromatin decondensation, Ag-NOR family, nucleolin antibody, systemic connective tissue disease, SLE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JNK1 antisense oligonucleotide, JNK2AS, (ISIS #12560).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 9 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                  Example 15; Col 41-42; 41pp; English.
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Best Local Similarity 88.9
Marches 16; Conservative
                                                                                                                                                                                                                                                                                                 Bennett CF, Cowsert LM;
                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                             WPI; 2001-079848/09.
                                                                                                                             Homo sapiens
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BXXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of Jun Kinase antisense mRNA for treating cancer by administering vector comprising promoter operably linked to DNA sequence that encodes the antisense mRNA to patient diagnosed with cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to the use of Jun Kinase (JNK) antisense oligomoucleotides for treating cancer and for screening compounds that minic or augment the effect of JNK antisense oligomorleotides treatment for cancer. The present sequence is one such JNK antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                /note= "This oligonucleotide is a 2'-0-methoxyethy] chimeric antisense oligonucleotide containing five MOED phoshopodiester residues flanking a 2'-deoxymucleotide/phosphorothioate region"
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      cancer;
phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
JNK; jun kinase; antisense; cytostatic; 2'-O-methoxyethyl oligonucleotide; MOE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Holbrook NJ;
                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                    /mod_base= OTHER
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Gaps

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Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03; 0; Mismatches 2; Indels

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immune response. The present sequence is one such immunostimulatory mucleic acide can be pyrimidine rich uncleic acide can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae) and/or critomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is slab useful for present sequence can also be used to redirect a minume deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                molecules encoding Mycoplasma hyppenumina Mip3 antigen. Mip3 antigen and its fragments are useful in manufacturing a vaccine for treating or preventing a disease or disorder in an animal, especially pig, caused by M. hyppenumoniae infection e.g. enzootic mycoplasma pneumonia. The mhp3-encoded proteins may be used as immunogens to generate antibodies which immunospecifically bind such an immunogen. The antibodies generated against the antigen are useful in diagnostic immunoassays, passive immunotherapy and generation of anti-idiotypic antibodies. Mip3 proteins may also be used in immunoassays, e.g. to detect or measure in a biological sample from a vaccinated or potentially infected test animal the presence of antibodies to the antigen, and thus to monitor the immune response and/or to diagnose infection of the animal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequence is an oligonucleotide primer used to sequence nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New apoprotein antigens encoded by mhp3 gene from Mycoplasma hyopneumoniae useful as a vaccine for treating or preventing diseases caused by Mycoplasma hyopneumoniae.
to a non-rodent subject in sufficient quantity to stimulate an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mycoplasma hyopneumoniae MHP3 antigen, sequencing primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MHP3; antigen; ss; vaccine; enzootic mycoplasma pneumonia; sequencing primer; antibody; immunoassay; immunotherapy;
                                                                                                                                                                                                                  present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                               Sequence 20 BP; 4 A; 7 C; 7 G; 2 Ț; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                        1678 GACTTCGGGCTGGCCCGG 1695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 27; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                  20 GACTTTGGCCTGGCCCGG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS03300 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mycoplasma hyopneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PFIZ ) PFIZER PROD INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequencing_primer; antil
anti-idiotypic antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-309781/33.
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AAS03300
$$$$$$$$$$$$$$$$$$
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                                                                                                                            Novel beta fructofuranosidase gene useful for producing beta fructofuranosidase and for developing variant enzymes that have increased heat resistance and transfer ratio by means of genetic engineering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vaccine, cytostatic, virucidal, bactericidal, fungicidal, anti-parasitic, immunostimulatory, tumour; viral infection; bacterial infection; fungal infection; parasitic infection, cancer, asthms, infectious disease; allergy, immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                 The present invention relates a beta-fructofuranosidase gene. The invetion is useful for the development of variant enzymes that have increased heat resistance and transfer ratio by means of genetic engineering techniques. Beta-fructofuranosidase is useful in the synthesis of transfructosylated oligosaccharides such as lactosucrose, for use in fields of foods and drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 72.2%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                     Ito T, Sakano Y, Fujita K;
            (NORQ ) SOC TECHNO-INNOVATION AGRIC FORESTY & FI.
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 5 C; 7 G; 1 T; 0 U; 4 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunostimulatory nucleic acid #299.
                                                                                                                                                                                                                                 Disclosure; Page 29; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           545
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
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27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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(COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CSGGSCCGTCCTGSAGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCGGCCCATCCTGCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF99183 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Krieg AM, Schetter C,
                                                     Tonozuka T,
                                                                                          WPI; 2001-123506/14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200122972-A2.
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                                                                                                                                                                                          techniques.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           528
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                                                   Hara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1620
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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic

diseases, allergies and asthma nucleic acids.

Vaccinating against tumors, infectious using immunostimulatory Py-rich and TG

WPI; 2001-273485/28.

Claim 101; Page 44; 338pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes isolated nucleic acids encoding secreted polypeptides, designated PROX polypeptides (i.e. a PRO polypeptide where X is an integer from 1 to 17). PROX polypeptides have cytostatic, immunomodulatory and reproduction activities, and can be used in gene therapy, and as PROX antagonists and PROX agonists. PROX polypeptides, nucleic acids and antibodies are useful in the manufacture of a medicament for treating a syndrome associated with a PROX-associated disorder, e.g. a cell proliferation and/or differentiation disorder (e.g. pre-clampsia). They are also used for screening for a modulator of activity or of latency or predisposition to a PROX-associated disorder. AAP74432 to AAP74448 encode the specifically claimed human PROX polypeptides PRO1 to PRO17 given in ABB70531 to AAB70547. The present
                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids encoding secreted polypeptides, designated PROX polypeptides, useful for treating a syndrome associated with a PROX-associated disorder, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                     Human, PRO; PROX; cytostatic; immunomodulatory; reproduction;
gene therapy; cell proliferation; differentiation disorder; cancer;
immune associated disorder; gestational disease; pre-clampsia;
                                                                  ö
                           Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                    Clone 7520500 PRO9 forward PCR primer SEQ ID NO:111.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other;
G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; Page 144; 166pp; English
                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer; sequencing primer; ss
                                                                                                 3539 GCTTCTAGAGTTTTATAG 3556
                                                                                                                                 19
                                                                                                                                                                                                                   AAF74514 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-AUG-1999; 99US-0148433P.
10-AUG-2000; 2000US-00635949.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-AUG-2000; 2000WO-US021857
                                             Local Similarity 88.9%;
les 16; Conservative
Sequence 20 BP; 3 A; 3 C; 5
                                                                                                                                 GCTTCTTCAGTTTTATAG
                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-147509/15.
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                                                                                                                                                                                                                                                                                     09-MAY-2001
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                                 Query Match
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                                                                  Matches
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The present invention describes a method of diagnosing age-related macular degeneration in a individual, involving comparing the level of eystatin C activity compared to a known reference value of a diseased or healthy individual. This is useful in identifying those at risk of and suffering from age-related macular degeneration. The cystatin C gene is found at human formomome 20pl.1.2 and the B allele has been shown to be indicative of an increased risk of the disease. The present sequence is PCR primer used to amplify the cystatin C gene (also known as CST3)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           involves comparing activity and/or level of transcriptional or translational product of Cystatin C gene in patient and control samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diagnosing or prognosing age-related macular degeneration, in a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense; E2F transcription factor 1; human; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                   Human; cystatin C; age-related macular degeneration; CST3; chromosome 20p11.2; polymorphism; B allele; PCR primer; Bs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                    Human cystatin C gene PCR primer #1206R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 30; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    (EVOT-) EVOTEC NEUROSCIENCES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2940 TGGAGGGAGGCCCCAGGG 2957
                 20
                                                                                            BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
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01-FEB-2000; 2000EP-00101921.
                                                                                                                                                                                                                                                                                                                                                         01-SEP-2000; 2000WO-EP008554
3 GGCCTGGACTGCTTCTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.48;
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                                                                                            AAF76793 standard; DNA; 20
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                                                                                                                                                         (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nitsch R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-235121/24.
                                                                                                                                                                                                                                                                                              WO200116364-A2
                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                        17-MAY-2001
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                                                                                                                                                                                                                                                                                                                            08-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Richard G,
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                                                                                                                         AAF76793;
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Gaps

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0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels

88.98;

Query Match
Best Local Similarity 88.9
Matches 16; Conservative

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FOR primers AAH75023-24 were used to amplify CDNA encoding human for fibroblast growth factor 23 (FGF-23). The human FGF-23 polymucleotide and colypeptide are useful for treating a patient suffering from traumatic injury or a condition characterized by dysfunction of or injury to skin cells, a condition characterized by inadequate function of placental condition characterized by inadequate function of placental condition characterized by inadequate function of the thymus (e.g. condition characterized by inadequate function of the thymus (e.g. condition characterized by inadequate function of the thymus (e.g. thymus, or differentiation disorder of the thymus), or a condition characterized by central nervous system disorder (e.g. Parkinson's condition characterized by central nervous system disorder (e.g. Parkinson's disease or Alzheimer's disease). The human FGF-23 polymucleotide and polypeptide are also useful in the treatment of Crohn's disease, healing confinemations of the intestine. They are also useful for treating canstomoses, motility and absorption disorders, and congenital consenses, motility and absorption disorders, and congenital consenses, motility and absorption disorders, stroke) and congenital consenses, renal artery disease, stroke) and congenting consenses are an expected or musculoskeletal disease characterized by loss of function, inadequate consenses, renal artery disease, stroke) and consenses of function or death of skeletal muscle cells, bone cells or supporting contracterized by loss of function, inadequate contracterized contract
                                                                                                                                                                factor 23, useful in the treatment of a condition characterized by inadequate function of placental cells (e.g. congenital defects) and the thymus (e.g. leukemia).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; systemic lupus erythematosus 1B; SLE-1B; dermatological; ss; antiinflammatory; immunosuppressive; systemic autoimmune disorder; signaling lymphocyte activation molecule; SLM», lymphocyte antigen 9; Ly-9; 2B4; natural killer cell receptor; CD48; CD84; LY108; CS1; DBDD; NIT1; upstream transcription factor 1; USF 1; GOLGA4; immune tolerance;
                                                                                                                                                         Isolated nucleic acids encoding the human and murine fibroblast growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human reverse PCR primer HMS1-26, used to amplify SlelB region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cells (e.g. skeletal myopathies, bone disease, or arthritis)
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88.9%; Pred. No. 1.7e+03;
iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                Disclosure; Page 8; 77pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             825 CTCTGCGTGGCTGGTGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                      Itoh N, Kavanaugh MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer; HMS1-26.
     (KYOU ) UNIV KYOTO.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to antisense compounds up to 30 nucleobases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pibroblast growth factor 23; FGF-23; injury; placental cell; ulcer; congenital defect; fertility; thymus; leukemia; lymphoma; injury; autoimmune disease; proliferative disorder; differentiation disorder; central nervous system disorder; parkinson's disease; inflammation; Alzheimer's disease; Crohn's disease; intestinal wound; stroke; motility disorder; absorption disorder; intestinal malformation; ischemic vascular disease; myocardial ischemia; myocardial infarction; peripheral vascular disease; renal artery disease; skeletal myopathy; musculoskeletal disease; skeletal muscle cell; bone disease; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in length targeted to a E2F transcription factor 1 The invention is useful for inhibiting the expression of E2F transcription factor 1 in cells or tissues. The antisense oligonucleotides may also be used as research agent and to prevent infection, inflammation or tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                Antisense compound capable of inhibiting the expression of E2F transcription factor 1, useful for preventing or delaying infection, inflammation or tumor formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer for human fibroblast growth factor 23 (FGF-23) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 0.4%; Score 14.8; DB 1; Length 20; Local Similarity 88.9%; Pred. No. 1.7e+03; les 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 4 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                     Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; Col 42; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18 AGAACCACATCCAGTGGC 1
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                                                                                                                                                 02-MAR-2000; 2000US-00517584
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18-SEP-2000; 2000US-023368P.
05-DEC-2000; 2000US-0251650P.
                                                                                                                                                                                                  02-MAR-2000; 2000US-00517584
                                                                                                                                                                                                                                                                                                  Popoff I, Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAR-2001; 2001WO-US007468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      620 AGCCCCACATCCAGTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH75023 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-190981/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200166595-A2
Homo sapiens
                                                 US6187587-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-OCT-2001
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                                                                                                  13-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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The present invention relates to a new method for screening for susceptibility to a systemic autoimmune disorder. The method comprises screening for at least one mutation within the systemic lupus erythematosus (SLE)-1B loci. Screening for susceptibility to autoimmune disorders such as systemic lupus erythematosus involves screening for at least one mutation in a gene or genes with the SLE-1B loci such as a gene encoding signalling lymphocyte activation molecule (SLAM), lymphocyte antigen (Ly)-9, 2B4 (a natural killer cell receptor), CD48, CD48, LY108, CSI, DEDD, NITI, upstream transcription factor (USF)1, GOLGA4. The method of the invention is useful for treating SLE and involves administering a mentioned genes. Gene therapy also involves the use of antisense constructs or ribozymes directed against the above mentioned genes for treating SLE. The present nucleic acid sequence represents the human
                                                                                                            Screening for susceptibility to systemic autoimmune disorder by screening for a mutation within the systemic lupus erythematosus-1B loci.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reverse PCR primer HMS1-26 that was used in the invention with the human forward PCR primer HMS1-26 (AAS20617) to amplify the human microsatellite marker HMS1-26 (AAS20619) for the S1elB region. HMS1-26 is also known as HMS1-25R and HMS1-4R and is located between the Ly-9 and the 2b4 genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Screening an unknown base sequence at a defined site of a target single-stranded nucleic acid for use in DNA diagnosis and therapy, comprises a DNA chip, fluorescence yield and pattern-based method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid base sequence analysis; DNA diagnosis; HSC5; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 3 A; 1 C; 9 G; 7 T; 0 U; 0 Other;
                                      Morel L;
                                                                                                                                                                  Example 6; Page 67; 128pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2325 GTGTGTGTGTGTGTGT 2342
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 88.9%;
Matches 16; Conservative
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                                    Wandstrat A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HSC5 exon 7 PCR primer E7A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-AUG-2002 (first entry)
 (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yamamoto N, Okamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-372310/40.
                                                                         WPI; 2002-066695/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CANO ) CANON KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200233068-A1.
                                    Wakeland EK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
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                                                  The present invention relates to a method of analysing an unknown nucleic acid base sequence. The method comprises preparing a probe array, hybridising with the probe array, measuring the fluorescence yield in the reaction, obtaining a template pattern, producing a sample pattern, and comparing the sample pattern with the template pattern. The method is useful for specifying an unknown base sequence at a defined site of a target single-stranded nucleic acid, which is useful for analysing a nucleic acid base sequence. The method is applicable in DNA diagnosis and therapy, and is useful in medicine and biology. Measuring the fluorescence yield allows the detection of a one-base mismatch which can be considered to produce high detection accuracy. The hybrid pattern of the DNA probe is used so the difference in themostability is less important, and the judgement on each spot can be reliably carried out. ABK72525-ABK72532 represent PCR primers used in the examples of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   administering at least one antiangiogenic nucleic acid molecule. Also
included is a kit comprising a first container housing the antiangiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to inhibiting angiogenesis in a subject, comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Angiogenesis inhibitor; 88; angiogenesis; 801id tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Angiogenesis inhibitory oligonucleotide #311.
                  4; Page 17; 53pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Page 25; 276pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2695 CCACTTCCCACCCTGCCC 2712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-DEC-2001; 2001WO-US048458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-DEC-2000; 2000US-025534P.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19 ccacrreccaccrecac
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                           present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bratzler RL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABS77827;
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                    Example
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cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, non-Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, pancreatic cancer, prostate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligomucleotide described in the exemplification of the invention

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0.4%; Score 14.8; DB 1; Length 20; 18.9%; Pred. No. 1.7e+03; Ive 0; Mismatches 2; Indels

88.98;

Conservative

Local Similarity es 16; Conserv

Best Loca Matches

Query Match

1678 GACTTCGGGCTGGCCCGG 1695

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20 cacriricacciadococa

BP

ABK49044 standard; DNA; 20

RESULT 1630

ABK4904

Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;

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          having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, theumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                         Antibody-induced cell lysis; cancer; immunostimulatory; CD20; anglogenesis; metastasis; cytostatic; phosphorothioate backbone; ss.
 and instructions for administering them to a subject
                                                                                                                                                                                                          .
0
                                                                                                                                                                              0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "phosphorothioate backbone"
                                                                                                                                                     Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                   Immunostimulatory nucleic acid SEQ ID NO: 463.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                 1678 GACTICGGGCTGGCCCGG 1695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER
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                                                                                                                                                                                                                                                                                                                        ВР
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                                                                                                                                                                                         88.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (IOWA ) UNIV IOWA RES FOUND.
                                                                                                                                                                                                                                                                                                                        ABL39057 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               base=
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                        Conservative
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/*tag=
                                                                                                                              acid of the invention
                                                                                                                                                                                          Local Similarity
 nucleic acids,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200197843-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
modified_base
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                                                                                                                                                                                                        16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                1629
                                                                                                                                                                                                     Matches
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Rice, fertility recovery gene, Rf-1 gene, RFLP marker, S12564, C1381, restriction fragment length polymorphism, chromosome 10, hybrid plant, rice BT type male fertility cytoplasm, G4003, PCR; primer, ss.

16-AUG-2001; 2001WO-JP007052. 17-AUG-2000; 2000JP-00247204.

WO200214506-A1.

Synthetic.

21-FEB-2002.

(NISB) JAPAN TOBACCO INC

(SYGN) SYNGENTA LTD

PCR primer #1, used to amplify G4003 HindIII marker sequence.

(first entry)

02-JUL-2002

ABK49044;

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The present invention relates to a new method of identifying rice individual or seed, for examination, which may contain a fertility eccovery gene (RF-1 gene). The method of the invention involves using the RF-1 gene locus located between the RFLP (restriction fragment length polymorphism) markers loci $12564 and Cl381 existing on the rice 10th chromosome. The method can be used for estimating the genotype of fertility recovery locus to rice piece stimating the genotype of certility recovery gene, which is useful in growing hybrid crice plants. Plural PCR marker loci located around and linked to the Rf-1 gene are developed and the relationship is defined. Therefore, the presence or absence of the Rf-1 gene can be examined, and an Rf-1 gene containtividual is selected conveniently and exactly by calibrating the genotypes of plural PCR marker loci. The present nucleic acid sequence represents PCR primar #1 that was used in the methods of the invention to amplify the G4003 HindIII marker sequence for use in detecting rice Rf-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Estimating genotype of fertility recovery locus to rice BT type mal fertility cytoplasm comprises detection of fertility recovery gene, useful in growing hybrid rice plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nitta N, Takemori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 31; 47pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene, as described above
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yamamoto T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-280759/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Komori T,
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The present invention relates to methods for treating or preventing devoloping administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx

Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of

Hartmann G;

Weiner G,

WPI; 2002-154611/20.

Disclosure; Page 212; 312pp; English.

developing cancer

PCR; primer; mhp3; apoprotein antigen; enzootic mycoplasmal pneumonia; vaccine; Mycoplasma hyopneumoniae infection; ss.

Mycoplasma hyopneumoniae

EP1245677-A1.

02-OCT-2002.

30-MAR-2001; 2001EP-00303030. 30-MAR-2001; 2001EP-00303030.

(PFIZ) PFIZER PROD INC. Madura RA,

King KW,

Mycoplasma hyopneumoniae mhp3 PCR primer #13.

16-JAN-2003

AAL50647;

AAL50647 standard; DNA; 20

RESULT 1632

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates in the primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the crealtant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis, AB142957 to AB145322 represent PCR primers for human chromosome 1986-35 DNA, and AB145323 to AB145634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                        Gaps
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
Length 20;
                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human chromosome 1p36-35 PCR primer SEQ ID NO:572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 2 C; 8 G; 7 T; 0 U; 0 Other;
Score 14.8; DB 1;
Pred. No. 1.7e+03;
                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 16; 528pp; Japanese.
                                                                                                                                                                                                                 2 Arceaceaeracereaac 19
                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-MAR-2000; 2000JP-00066716.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAR-2001; 2001JP-00068285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                         2044 ACCGACGAGTACCTGGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                    ABL43528 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JP2001321190-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-NOV-2001.
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Best Local S
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                                                                                                                                                                                                                                                                                                                            RESULT 1631
ABL43528/c
XX
AC ABL4352
XX
DT 11-APR-
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Human;
XW
Human;
XW
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                                                                                                                                                                                                                                                                                                                                                                                                     Novel apoprotein antigens encoded by Mycoplasma hyopneumoniae for use in vaccines to prevent and treat diseases caused by infection with Mycoplasma hyopneumoniae in animals, especially pigs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Appropriate antigens encoded by the M. hyopneumoniae mip3 gene. M. hyopneumoniae is a bacterial pathogen that causes enzootic mycoplasmal pneumonia in pigs. The mip3 genes, proteins and apoprotein antigens of the invention are useful in the manufacture of a vaccine for treating/ preventing a disease or disorder caused by infection with M. hyopneumoniae in an animal, especially a pig. The present DNA sequence represents a PCR primer for the Mycoplasma hyopneumoniae mip3 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Candida albicans GRACE strain PCR primer SEQ ID NO 4779.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 3 C; 5 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3539 GCTTCTAGAGTTTTATAG 3556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 26; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCTTCTTCAGTTTTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABZ30628 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-742716/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ30628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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ABZ30628/c
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Gaps

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999 CCCCACCGTGCACAAGAT 20 CCACACCATGCACAGAT

16; Conservative

Matches

Local Similarity

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sapiens.
                   Synthetic.
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Matches
Homo
                                                           Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cone allele by insertion or replacement by a cassette having an expressible selectable marker and modifying other allele by resolution, of a promoter replacement fragment with a heterologous promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles modified are modified. The diploid fungal cells in which both alleles modified are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that is essential to the survival or growth of a fungus, a gene that contributes to the virulene and/or pathogenicity of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent, an antifungal agent that inhibits the growth of a diploid fungus or and for identifying a therapeutic agent for treatment of a mammalian disease. (M1) is useful for identifying a compound which modulates the activity of a gene product, preferably enzymatic activity, carbon compound catabolism, blosynthetic, transporter, transcriptional, compound catabolism, blosynthetic, transporter, transcriptional, activity of inhibit growth or proliferation of C. albicans cells and for treating infection by C. albicans. The present sequence is that of a PCR primer used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based to nequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                            Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; raf; hyperproliferation; neovascularisation; ocular angiogenesis; therapy; cancer; cytostatic; anti-angiogenic; vascular; ophthalmological; antisense; phosphorothioate backbone; c-raf kinase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 36; SEQ ID NO 4779; 167pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human c-raf kinase antisense oligonucleotide ISIS #7854.
                                                                                                                                                                                                                                                               Ohlsen KL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 7 A; 11 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                               Bussey H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          846
                                                                                                                                                                                                                                                               Boone C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19 dreferciedrecie 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD44739 standard; DNA; 20 BP.
                                                                                                                                        29-DEC-2000; 2000US-0259128P
                                                                                                                                                           20-FEB-2001; 2001US-00792024
22-AUG-2001; 2001US-0314050P
                                                                                               26-DEC-2001; 2001WO-US049486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            829 GCGTGGCTGGTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                       (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 88.5 es 16; Conservative
                                                                                                                                                                                                                                                               Roemer T, Jiang B,
                                                                                                                                                                                                                                                                                                      WPI; 2002-566694/60
                 WO200253728-A2
                                                       11-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1634
AAD44739/c
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capable of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to novel antisense oligonucleotides which are targetted to nucleic acids encoding human raf proteins and capable cinhibiting raf expression. The invention also relates to methods of inhibiting hyperproliferation of cells which involves contacting the hyperproliferation of cells which involves contacting the hyperproliferation of the invention. The method is useful for treating cancer, angiogenesis or neovascularisation, especially ocular angiogenesis or neovascularisation. The present DNA sequence is an antisense oligonucleotide targetted to human c-raf kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Phosphorothioate; HIV-1; azasugar; AIDS; virucide; antiviral; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating cancer, angiogenesis or neovascularization by administering antisense oligonucleotides targeted to human raf sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
                                                                                                  'note= "Phosphorothioate backbone'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Phosphorothioate oligonucleotide for AIDS therapy.
                                                                                                                                                                              /mod_base= OTHER
/note= "2'-0-methyl nucleotides"
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure, Col 14; 41pp; English
                                                                            /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  844 CTGCCAGCCGAGGAG 861
                                                                                                                                                                                                                                                                                                                                                                                                                                            95WO-USO07111.
96US-00756806.
97US-00888982.
98WO-US013961.
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                                                                                                                                                                                                                                                                                                                                                                     2000US-00506073
                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00250856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-00143214.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 88.9
nes 16; Conservative
                           ..20
*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-597918/64.
                        modified_base
                                                                                                                             nodified base
                                                                                                                                                                                                                                                                                                                                                                     18-FEB-2000;
                                                                                                                                                                                                                                                             US6410518-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUL-1997;
                                                                                                                                                                                                                                                                                                                 25-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monia BP
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Human, K-ras, PCR primer; probe, capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                      Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                                                                                                                                                                             (CORR ) CORNELL RES FOUND INC
                                                                                                                                                             04-APR-2001; 2001WO-US010958.
                                                                                                                                                                                      14-APR-2000; 2000US-0197271P.
                                                                                                                                                                                                                                      Barany F, Zirvi M,
                                                                                                                                                                                                                                                               WPI; 2002-034366/04.
                                                                                                            WO200179548-A2.
                                                                                                                                     25-OCT-2001
                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABI96621;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is that of a phosphorothioate oligonuclectide of random sequence which includes 4 six-membered azasugar nucleotide derivatives. It is a claimed example of oligonucleotides of the invention (see ABV73816-41) that have been tested as AIDS therapeutic agents. In anti-HIV-1 assays, the oligonucleotide showed higher antiviral activity than AZT, ddC and ddI, and antiviral activity was resistant to the effects of serum. Claimed oligonucleotides of the present invention have low toxicity against cells, are membrane permeable, working outside of cells to inhibit viral attachment of HIV, have a wide antiviral activity against a broad spectrum of HIV variants, are not active against other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present oligonucleotide to
                                                                                                                                                                                                                                                                                                                                                                                                                                                        New phosphorothioate oligonucleotides useful in the treatment of AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                  'note= "azasugar-containing adenosine derivative"
                                                                                                                                                                                      /mod_base= OTHER
/note= "azasugar-containing adenosine derivative"
                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "azasugar-containing adenosine derivative"
                                                                                                                                                note= "azasugar-containing adenosine derivative"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                serum allows its use as an AIDS therapeutic drug in vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 9 C; 4 G; 3 T; 0 U; 0 Other;
                                                /note= "phosphorothioate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                       Jung K;
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                                                                                                                                                                                                                                                                                                                                                                                                       Lee K,
Location/Qualifiers
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/mod base= OTHER
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                                    /mod base= OTHER
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Best Local Similarity 88.99
Watches 16; Conservative
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Key
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Favis R, Kliman R;

Gerry NP,

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(I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and functious agents e.g. Cryptococcus neoformans, Candida albicans and Sapergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, pepteinin-Bar virus and polio virus, and parasitic infectious agents selected from Onchovera volvulus, Entamoba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and Obselty defects.

Detecting cancer involving oncogenes, thuncr suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the electron accorded and correlating (using a computer) identified problem and correlating (using a computer) identified brook and correlating (using a computer) identified brook and correlating comprises computers) identified brook and correlating comprises computers) identified brook and correlating computers in computers in the angelon of the oligonucleotide probesed and correlating computers.
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ABI97546 represent oligonucleotide sequences used in the exemplification
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                                                                                      The present invention describes a method (MI) for designing
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38.9%; Pred. No. 1.7e+03;
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Example 5; Fig 29; 300pp; English
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Matches 16, Conservative
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ID AB19
XX AC AB10
XX DT 16-
XX DE CAP
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Capture oligonucleptide Zip ID#845 oligo #9.

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oligonucleotide probes (II) will hybridise with little mismatch, where oligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious agents e.g. caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Sapergillus funigatuts, viruses e.g. T-cell lymphocytotrophis cirus, Espetein-Barr virus and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCAI gene, p53 gene, human papillomavirus types I6 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the contract and correlating (using a computer) identified brook as a set in a computer of the oligonucleotide probe and infrared microscope) the support at the
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                       Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCAl; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obsity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
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ABI95745/C
ID ABI95745
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DT Capture
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The present invention describes a method (MI) for designing capture oligonucleotide probes (I) for use on a support to which complementary coligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detection infectious diseases caused by bacterial infectious agents (C) for detectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigatus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents or estected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC selected from Onchoverva volvulus, interacting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, buman papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe sets occurred and correlating (using e computer) identified ligation to a presence or absence of the target uncleotide sequences. Balls2074 to fereses the support and infrared microscope and incleotide sequences used in the exemplification
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                     Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
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Pred, No. 1.7e+03;
0; Mismatches 2; Indels
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Best Local Similarity
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The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) will hybridise with little mismatch, where for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Cadida albicans and Aspergillus funigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents as elected from Onchoverava volvulus, Entemmeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC Detecting cancer involving oncogenes, tumour suppressor genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, buman papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probeset sets and correlating (using a computer) identified ligation to a presence or absence of the target uncleotide sequences. ABIRB2074 to a base of the target uncleotide sequences used in the exemplification
                   Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
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                                                                                                                                                                                                                                                                                                      04-APR-2001; 2001WO-US010958.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 88.9
Marches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                          Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-034366/04.
                                                                                                                                                                                                        WO200179548-A2
                                                                                                                                                                                                                                                        25-OCT-2001,
                                                                                                                                                                                                                                                                                                                                                                                                                                             Barany F,
                                                                                                                                                          Synthetic
셤
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RESULT 1641
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                                Gaps
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0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                ABI97200 standard; DNA; 20 BP.
                                                                                         2 GCGGGTACTGTGCAACTT
                                                                                                                                                                                               ABI97200;
                                                                                                                                      RESULT 1640
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Capture oligonucleptide Zip ID#4287 oligo #9.

(first entry)

16-FEB-2002

SXXXXXX

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The present invention describes a method (MI) for designing deprute of conground-sociale probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents c.g. Cryptococcus neoformans, Candida albicans and infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus funigatuts, viruses e.g. T-cell lymphocytotrophis cirus, Estected from Onchoverva volvulus, Entamoeba histolycica and bracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC betecting cancer involving oncogenes, tumour suppressor genes or involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, cancer is superioration for environmental monitoring (using e.g. a scanning cand feed industry, detecting comprises scanning (using e.g. a scanning calcutulated and infrared microscope) the support at the electron microscope and infrared microscope) the support at the cancer sets sets of the target nucleotide sequences or absence of the target nucleotide sequences or absence of the target nucleotide sequences or absence of the target nucleotide sequences. Appression to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AB197546 represent oligonucleotide sequences used in the exemplification
                   Human, K-ras, PCR primer; probe, capture probe, mutation detection; ligase detection reaction; LNR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rice chromosome 10 RFLP marker G4003 HindIII PCR primer, SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a method (M1) for designing capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gerry NP, Favis R, Kliman R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Fig 29; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1692 CCGGGACGTGCACACCT 1709
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                                                                                                                                                                                                                                                                                                                                                                                        (CORR ) CORNELL RES FOUND INC.
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                                                                                                                                                                                                                                                                                                                                             14-APR-2000; 2000US-0197271P.
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                      Barany F, Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-034366/04
                                                                                                                                                                                                    WO200179548-A2.
                                                                                                                                                                                                                                                    25-OCT-2001.
                                                                                                                                                            Synthetic.
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ID ACF3
XX
AC ACF3
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DT 22-E
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DE RICE
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real time quantitative PCR; RTQ PCR; primer; ss.
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 The invention relates to the restoration to and control of fertility in transforming rate with BT cytoplasmic male sterility. The invention involves transforming rate with a nucleic acid containing all or part of the fertility restorer gene (Rf-1) derived from rice cultivar IR24 (ACF30938), or a sequence at least 70% homologous to it which encodes a protein with similar activity. The Rf-1 gene is located on chromosome 10, and was analysed using RFUP (restriction fragment length polymorphism) and PCR to determine markers. The Rf-1 gene product is thought to be responsible for reducing the abnormal transcription of the mitochondrially-encoded atp6 gene which is thought to be responsible for cytoplasmic male sterility. The invention also encompasses a method of determining the presence of the Rf-1 gene in a rice genome, using suitable primers to amplify from the rice genome a marker consisting of the P4497 Mbo1-B56691 XbaI fragment of rice chromosome 10, and nucleic acids containing residues 3838-54123 or 42132-48883, and homologous sequences may be used to control and/or restore fertility in rice having BT cytoplasmic male sterility. Sequences ACF30912-ACF30929 and ACF30940-ACF30979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                  Fragment of rice Rf-1 gene for restoration to and control of fertility in rice with BT male cytoplasmic sterility.
           Rice; plant; fertility; control; restoration; cytoplasmic male sterility; BT; fertility restorer; Rf-1; chromosome 10; marker; RFLP; restriction fragment length polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; NOVX; G-protein coupled receptor; GPCR; cancer; cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer #2 for DNA encoding human NOV19a.
                                                                                                                                                                                                                                                 Komori T, Ota S, Murai N, Hiei Y;
                                                                                                                                                                                                                                                                                                                                        Example; Page 96; 200pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2044 ACCGACGAGTACCTGGAC 2061
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04-OCT-2001; 2001JP-00309135.
26-JUN-2002; 2002JP-00185709.
                                                                                                                                     13-SEP-2002; 2002WO-JP009429.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABX70535 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-MAR-2003 (first entry)
                                                                                                                                                                                                             (NISB ) JAPAN TOBACCO INC (SYGN ) SYNGENTA LTD.
                                                                                                                                                                                                                                                                            WPI; 2003-313641/30.
                                                                                   WO2003027290-A1
                                                             Oryza sativa
                                                                                                            03-APR-2003.
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Kekuda R, Tchernev VT, Liu X, Spytek KA, Patturajan M;
Burgess CE, Vernet CAM, Li L, Gorman L, Malyankar UM, Boldog FL;
Guo X, Shenoy SG, Padigaru M, Taupier RJ, Miller CE, Camman SJ;
Pena CEA, Gangolli EA, Gusev V, Smithson G, Zerhusen BD, Gerlach V;
Pochart PF, Fernandes ER, Shinkets RA, Rastelli L, Spaderna SK;
Larochelle WJ, Zhong M, Khramtsov NV, Voss EZ, Herrmann JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NOVX polypeptides and polynucleotides, useful for treating a syndrome related to a human disease associated with the NOVX polypeptide e.g.,
                                                                                                                                                                                                                                                                                        2001US-0275579P.
2001US-0275601P.
2001US-0276000P.
2001US-027676P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001US-0277338P.
2001US-0277791P.
2001US-0277833P.
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2001US-0278999P.
2001US-0279036P.
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2001US-0310913P.
2001US-0311978P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2001; 2001US-vcccc
02-MAY-2001; 2001US-0288228P.
17-MAY-2001; 2001US-0291766P.
                                                                                                                                   08-MAR-2002; 2002WO-US007355.
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2001US-0277327P.
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2001US-0318403P.
2001US-0318510P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-0313626P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001US-0322296P.
2001US-0322360P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001US-0280233P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001US-0314018P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-058423/05.
                                                                                                                                                                                                                                                                     13-MAR-2001; 2
13-MAR-2001; 2
13-MAR-2001; 2
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAR-2001;
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21-AUG-2001;
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                                                                                       10-OCT-2002
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with RNA or DNA, comprising covalently-bound nucleosides that individually include a ribose of deoxyribose sugar portion and a base portion. The nuclease resistant compounds are useful as therapeutics, diagnostic agents, or research reagents. The compounds are also useful for modulating the activity of an RNA or DNA molecule, or for treating ar organism with a disease associated with the undesired production of a protein, e.g. bacterial, viral or protozona infections, AIDS, abnormal cell proliferation and tumour formation, or atherosclerosis. The present sequence represents the human c-raf mRNA antisense oligonucleotide #5
                                           The invention relates to a nuclease resistant compound that hybridises
                                                                                                                                                                                       Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                   Example 31; Page 29; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200302227-A2
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                                                                                                                                                                                                                     RESULT 1644
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      $$$6666666668$$
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The present invention relates to the isolation of novel human polypeptides referred to as NOVX (NOVI-NOV44), variants of these proteins, and the polymucleotide sequences encoding them. The NOVX proteins of the invention are G-protein coupled receptor (GPCR) related proteins. The sequences of the invention are useful in the manufacture of a medicament for treating a syndrome related to a human disease associated with the polypeptides e.g. cancer. The present sequence represents a PCR primer used in a real time quantitative (RTQ) PCR reaction for DNA encoding a human NOVX protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nuclease resistant compounds, useful as therapeutics, diagnostic agents, or research reagents, or for treating an organism with a disease associated with the undesired production of a protein, e.g. bacterial infections or AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .20
/*tag= a
/mod_base= OTHER
/note= "OTHER = phosphorothioate backbone. Optionally 1-
12 are 2'-O-methyl nucleotides"
                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                Human, c-raf, antisense, ss; nuclease inhibitor; gene therapy; AIDS;
bacterial infection; viral infection; protozoan infection;
abnormal cell proliferation; tumour formation; atherosclerosis.
                                                                                                                                                                                         ö
                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
                                                                                                                                     Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                        Human c-raf mRNA antisense oligonucleotide #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                2008 GTGGAGGACCTGGACCGT 2025
                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90US-00463358.
90US-00566977.
91WO-US000243.
91WO-US005720.
91US-00814961.
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92US-00854634.
92WO-US011339.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94US-00244993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-00471973
98US-00135202
                                                                                                                                                                                                                              3 GAGGAGGACCTGGACAGT
                                                                                                                                                                                                                                                                                                         ACA61357 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cook PD, Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-438873/41.
                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2003004325-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified base
                                                                                                                                                                                                                                                                                                                                                              11-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-JAN-1991;
12-AUG-1991;
24-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-JAN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                     ACA61357;
                                                                                                                                                                                                                                                                                  RESULT 1643
                                                                                                                                                                                        Matches
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/note= "This oligonucleotide has a phosphorothioate
/note= "This oligonucleotide has a most wings at the 5'
backbone and 2'-O-methoxyethyl (2'-MOE) wings at the 5'
and 3' ends, which are 5 nucleotides in length. Also all
cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotide targeted to a nucleic acid encoding vascular endothelial growth factor receptor-1, useful for diagnosing or treating cancer, rheumatoid arthritis, or diseases or conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vascular endothelial growth factor receptor 1; VEGF receptor; VEGFR; inhibitor; cytostatic; antirheumatic; antiarthritic; antiangiogenic; antibitory; antisense quencherapy; hyperproliferative disorder; cancer; rheumatoid arthritis; angiogenesis; infection; inflammation; tumour formation; phosphorothioate; 2'-O-methoxyethyl; 2'-MOE; ss.
                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human VEGFR-1 chimeric phosphorothioate oligonucleotide SEQ ID NO:97
                                                             ö
Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
                                                          2; Indels
                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                     844 CTGCCAGCCGAGGAGGAG 861
                                                                                                                                                                             m
                                                                                                                                                                                                                                                                                                                                     ВР
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0.4%;
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                                                                                                                                                                             20 creccaegesaceae
                                                                                                                                                                                                                                                                                                                                  ACC86802 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ಹ
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   Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     involving angiogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennett CF, Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-301004/29.
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The present interaction describes a compound (v) of our currecoper in the present interaction described encoding vascular endothelial growth factor receptor-1 (VEGFR-1), where the compound inhibits the expression of VEGFR-1 or with an 8-mucleobase portion of an active site on the nucleic acid molecule encoding VEGFR-1. Also described: (1) a composition of acid molecule encoding VEGFR-1. Also described: (1) a composition of comprising (C) and a carrier or diluent; (2) inhibiting the expression of VEGFR-1 in cells or tissues by contacting the cells or tissues with (C) or that the expression of VEGFR-1 is inhibited; and (3) treating an animal so that the expression of VEGFR-1 is inhibited. (C) has antianglogenic, antirheumatic, antiarthritic, administering (C) to the animal so that the expression of VEGFR-1 is inhibited. (C) has antianglogenic, antirheumatic, antiarthritic, cytostatic and antiinflammatcopy activities, and can be used in antisense compounds are used in antisense compounds compounds are used in antisense (e.g. cancer), rheumatoid arthritis, or diseases or conditions associated with the expression of VEGFR-1, such as hyperproliferative disorders (c) e.g. cancer), rheumatoid arthritis, or diseases or conditions involving anglogenesis. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, confitcing the temper for the entire members of a biological distinguishing between functions of various members of a biological prophylaxis, e.g. to prevent or delay infection.

Company The present sequence represents a human VEGFR-2 chimeric pathway. The present inventions
                                    present invention describes a compound (C) 8-50 nucleobases in length
X3333333333333333333333333333333X8
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0.4%; Score 14.8; DB 1; Length 20; 18.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels Sequence 20 BP; 5 A; 2 C; 8 G; 5 T; 0 U; 0 Other; 1352 TGGAGATGATGAAGATGA 1369 88.98; 16; Conservative Best Local Similarity Query Match Matches à 셤

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Gaps

. 0

Human JNK2 sense control oligonucleotide ISIS12560. ADA26589 standard; DNA; 20 BP (first entry) 20-NOV-2003 ADA26589; RESULT 1645

88; human; Jun N-terminal kinase; JNK1; JNK2; JNK3; cytostatic; antiinflammatory, apoptosis, prostate cancer, prostate tumour, inflammation, fibrosis; fibrosic disease; fibrotic scarring, peritoneal adhesion, lung fibrosis; conjunctival scarring, hyperproliferative disease, cancer; probe.

Homo sapiens,

JS2003004120-A1. 02-JAN-2003.

31-JAN-2001; 2001US-00774809

97US-00910629. 98US-00130616. 99US-00287796. 99US-00396902. 13-AUG-1997; 07-APR-1999; 15-SEP-1999; 07-AUG-1998;

NERO P. GAARDE W A. (MCKA/) MCKAY R. (DEAN/) DEAN N M. (MONI/) MONIA B P. (NERO/) (GAAR/)

Gaarde WA; Nero P, Monia BP, Dean NM, Mckay R,

WPI; 2003-311908/30

the expression of condition New oligonucleotides which hybridizes to, and modulates the expr Jun N-terminal kinase, useful for treating a disease or conditio characterized by a reduction in apoptosis, e.g. prostate cancer, inflammation or fibrosis.

Example 4; Page 26; 69pp; English.

The invention relates to an oligonuclectide (antisense, AS) comprising 830 nuclectides connected by covalent linkages, where the oligonuclectide
30 nuclectides connected by covalent linkages, where the oligonuclectide
30 nuclectides specifically hybridisable with a nucleic acid encoding a
30 Jun N-terminal kinase (JNK) protein and modulates the expression of the
30 Jun N-terminal kinase (JNK) protein and modulates the expression of the AS oligonuclectide (or its bioequivalent, and a pharmaceutical
30 Treating an animal having/suspected of having/prone to having a
31 Carrier), treating an animal having/suspected of having/prone to having a
32 Carrier of the composition of the AS oligonuclectide),
33 Contacting the calls or tissues with the AS oligonuclectide, modulating
34 Contacting the calls or tissues with the AS oligonuclectide, modulating
35 Contacting the calls or tissues with the AS oligonuclectide, modulating
36 Contacting the calls or tissues with the AS oligonuclectide, modulating
37 Contacting the calls or the phosphorylation of a callular protein
38 Contacting the calls or the phosphorylation of a callular protein
39 Conduction and animal by administering the oligonuclectide to the calls of the conduction associated with a JNK protein or characterised by a reduction
39 Conduction associated with a JNK protein or characterised by a reduction
30 Conduction associated with a JNK protein or characterised by a reduction
30 Conduction associated with a JNK protein or characterised by a reduction
30 Conduction associated with a JNK protein or characterised by a reduction
30 Conduction with a JNK protein or characterised by a reduction
30 Conduction (e.g. fibrotic scarring), peritoneal adhesions, lung
31 Conduction, such as cancer. The antisense oligonuclectides may also be
31 Conduction, such as cancer. The antisense oligonuclectides may also be
32 Conduction, such as cancer. The mutisense oligonuclectides may also be
33 Conduction of the function of ne or near early or the presence or condition of ne or s function of one or more genes in the animal. The present is a sense control oligonucleotide for antisense oligonucleotides targeting a human JNK. sequence

Sequence 20 BP; 2 A; 7 C; 7 G; 4 T; 0 U; 0 Other;

ö 0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03; 2; Indels 0; Mismatches 88.9%; 16; Conservative Local Similarity Query Match Matches

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Gaps

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ADA26578 standard; DNA; 20 BP ADA26578; RESULT 1646 ADA26578,

Human Jun N-terminal kinase, JNK2, antisense oligonucleotide ISIS12560. ss; human; Jun N-terminal kinase; JNK1; JNK2; JNK3; antisense;

(first entry)

20-NOV-2003

cytostatic; antiinflammatory; apoptosis; prostate cancer; prostete tumour; inflammation; fibrosis; fibrotic disease; fibrotic scarring; peritoneal adhesion; lung fibrosis; conjunctival scarring; hyperproliferative disease; cancer; probe.

Homo sapiens

Location/Qualifiers modified base Key

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The invention relates to an oligonucleotide (antisense, AS) comprising 8-
30 nucleotides connected by covalent linkages, where the oligonucleotide
30 nucleotides connected by covalent linkages, where the oligonucleotide
30 nucleotides geotifically hybridisable with a nucleic acid encoding a
30 Jun N-terminal kinase (JMK) protein and modulates the expression of the
30 oligonucleotide (or its bioequivalent, and a pharmaceutical
31 carrier) treating an animal having/suspected of having/prone to having a
32 carrier) treating an animal having/suspected of having/prone to having a
33 chyperproliferative disease (by administering to a prophylactic or
34 therapeutic amount of the composition of the AS oligonucleotide,
35 contacting the cells or tissues with the AS oligonucleotide, modulating
36 the cells or tissues with the AS oligonucleotide, modulating
37 the cell cycle progression (or the phosphorylation of a protein
38 contacting the cells or the phosphorylation of a protein
39 promotes one or more metaatatic events in cultured cells or the cells of
30 the growth of a tumour in an animal by administering the oligonucleotide to the cells, inhibiting
39 trowncleotide for JNK2 and treating a cell with an AS
30 condition associated with a JNK protein or characterised by a reduction
30 in apoptosis by administering a prophylactic or therapeutic amount of the
30 oligonucleotide. The animal subsection in apoptosis, such as
30 condition associated with a JNK protein or therapeutic amount of the
30 condition associated unmur, inflammation, fibrosis or condition (e.g. fibrotic scarring, peritoneal adhesions, lung
31 fibrosis or conduction (e.g. fibrotic scarring, peritoneal adhesions, lung
32 fibrosis or conduction (e.g. fibrotic scarring, peritoneal adhesions, lung
33 conduction, such as cancer. The antisense oligonucleotide may also be
34 cused as research agents and diagnostic aids, to detect the presence
35 conduction of one or more genes in the animal. The presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotides which hybridizes to, and modulates the expression of Jun N-terminal Kinase, useful for treating a disease or condition characterized by a reduction in apoptosis, e.g. prostate cancer, inflammation or fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence is an antisense oligonucleotide targeting human JNK2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03;
                    /mod_base= OTHER
/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaarde WA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 25; Page 25; 69pp; English.
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                                                                                                                                                                                                                                         98US-00130616
99US-00287796
                                                                                                                                                                         2001US-00774809
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                                                                                                                                                                                                                                                                                                                                MCKAY R.
DEAN N M.
MONIA B P.
                                                                                                                                                                                                                                                                                                                                                                                                NERO P. GAARDE W A.
                                                                                      US2003004120-A1.
                                                                                                                                                                           31-JAN-2001;
                                                                                                                                                                                                                                                             07-APR-1999;
15-SEP-1999;
                                                                                                                                02-JAN-2003
                                                                                                                                                                                                                                             07-AUG-1998
                                                                                                                                                                                                                                                                                                                                (MCKA/) 1
(DEAN/) 1
(MONI/) 1
(NERO/) 1
(GAAR/) 0
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                                                                                                                                                       !ransgenic; nucleoprotein; recombinase; lysozyme; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method for producing transgenic livestock animal e.g. pig, by introducing nucleoprotein made of nucleic acid and recombinase into totipotent or pluripotent cell, and growing the resulting recombinant totipotent or pluripotent cell. The method is useful for producing transgenic livestock animal such as pigs, goats, sheep, cows or horses, preferably goats and pigs. The present sequence is a primer used for amplifying human lysozyme cDNA. This sequence is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  congenital heart defect; prostate cancer; diabetes; metabolic disorder; neoplasm; graft versus host disease; AIDS; bronchial asthma; primer; crohn; a disease; multiple sclerosis; infectious disease; anorexia; cancer-associated cachexia; neurodegenerative disorder; RT-PCR; Alzheimer's disease; Parkinson's disease; immune disorder; haematopoietic disorder; dyslipidaemia; wasting disorder; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing transgenic livestock animal e.g. pig, by introducing nucleoprotein made of nucleic acid and recombinase into totipotent or pluripotent cell, and growing the resulting recombinant totipotent or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; 88; PCR; NOVX; cardiomyopathy; atherosclerosis; hypertension;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 20;
8.9%; Pred. No. 1.7e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                 Anderson GB, Murray JD, Oppenheim SM;
                                                                                                                    Human lysozyme cDNA amplifying primer, HL4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Forward RT-PCR primer for human NOV3e/f.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 30; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      386 TCAAGCTGCGGCATCAGC 403
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                AAD55160 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                              07-SEP-2001; 2001US-0317915P.
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                                                                                  (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-313182/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pluripotent cell.
                                                                                                                                                                                                                         WO2003022220-A2.
                                                                                                                                                                                           Homo sapiens.
                                                                                  07-AUG-2003
                                                                                                                                                                                                                                                           20-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                  Maga EA,
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                                                AAD55160;
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AAD55160
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Gaps ö

2; Indels

0; Mismatches

1678 GACTTCGGGCTGGCCCGG 1695

16; Conservative

Best Loc Matches

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20 GACTTTGGCCTGGCCCGG

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The invention relates to an isolated polypeptide comprising one of 127 sequences (appearing as ABO1288-ABO1414) designated as NOVX, a mature form of NOVX, an amino acid sequence comprising one or more conservative NOVX or an amino acid sequence comprising one or more conservative proteins, determining the presence or amount of NOVX or NOVX DNA in a sample (by introducing the sample to an antibody that binds immunospecifically to the polypeptide, and determining the presence or amount of antibody bound to the polypeptide, determining the presence of or predisposition to a disease associated with altered levels of expression of NOVX or NOVX DNA in a first mammalian subject, identifying a spectation to a disease associated with altered levels of expression of NOVX or NOVX identifying a potential therapeutic agent the physiological interactions of NOVX, screening for a modulator of activity of or of latency or predisposition to a pathology associated with NOVX, a vector comprising NOVX DNA, a cell comprising the vector (used to produce NOVX) and an anti-NOVX antibody. The NOVX nucleic acids and polypeptides are useful as a marker for cell or tissue type, and in diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polypeptides and polynucleotides having properties related to stimulation of biochemical or physiological responses in a cell or tissue, useful for diagnosing or preventing e.g. atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhong M, Li L, Gorman L, Spytek KA, Kekuda R, Taupier KJ; Anderson DW, Vernet CAM, Catterton B, Miller CE, Shenoy SG; Patturajan M, Pena CEA, Tchernev VT, Padigaru M, Gusev VY; Malyankar UM, Burgess CE, Gerlach VL, Casman SJ, Rieger DK; Grosse WM, Smithson G, Peyman JA, Starling G, Rothenberg ME; Larochelle WJ, Shimkets RA, Crabtree J, Rastelli L, Voss EZ; Boldog FL, Edinger SR, Millet I, Macdougall JR, Ellerman K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example C; Page 498; 849pp; English.
                                                                                                                                                                                                                                          12-SEP-2001; 20010S-0118450F.
17-SEP-2001; 2001US-0318765P.
17-SEP-2001; 2001US-032281EP.
19-SEP-2001; 2001US-0323819F.
20-SEP-2001; 2001US-0323631P.
25-SEP-2001; 2001US-032366P.
25-SEP-2001; 2001US-0324999P.
25-SEP-2001; 2001US-0324999P.
15-FEB-2002; 2001US-0324999P.
15-FEB-2002; 2001US-0324999P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002US-0366131P.
2002US-0367753P.
2002US-0369479P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypertension, prostate cancer
                                                                                                                                                                                                                2001US-0318130P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-MAY-2002; 2002US-0379532P.
17-MAY-2002; 2002US-0381664P.
17-MAY-2002; 2002US-0381672P.
28-MAY-2002; 2002US-0383651P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUN-2002; 2002US-0390155P.
                                                                                                                                                     09-SEP-2002; 2002WO-US028596
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   reverse transcriptase PCR
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                                                                            WO2003023008-A2.
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25-MAR-2002; 2
02-APR-2002; 2
                                           Homo sapiens.
                                                                                                                                                                                                              07-SEP-2001;
10-SEP-2001;
                                                                                                                  20-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chapoval A;
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NOVX sequences, including cardiomyopathy, atherosclerosis, hypertension, congenital heart defects, prostate cancer, diabetes, metabolic disorders, neoplasm, graft versus host disease, AIDS, bronchial asthma, Crohn's disease, multiple sclerosis, infectious diseases, anorexia, cancerassociated cachexia, neurodegenerative disorders (e.g. Alzheimer's disease or Parkinson's disease), immune disorders, haematopoietic disease or Parkinson's disease), immune disorders, haematopoietic diseases. These may also be used to screen for molecules which inhibit or enhance NOVX activity or function, and for detecting specific cell types. These may also be used in chromosome mapping, gene therapy, tissue typing, and in forensic biology. The present sequence is a reverse transcriptase (RT)-PCR primer used to assess the tissue specific expression of mRNA encoding a NOVX protein
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   conditions or disorders associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  congenital heart defect; prostate cancer; diabetes; metabolic disorder; neoplasm; graft versus host disease, ALDS; bronchial asthma; primer; crohn's disease; multiple sclerosis; infectious disease, anorexia; cancer-associated cachexia; neurodegenerative disorder; RT-PCR; Alzheimer's disease; Parkinson's disease; immune disorder; RT-PCR; haematopoietic disorder; dyslipidaemia; wasting disorder; gene therapy; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; ss; PCR; NOVX; cardiomyopathy; atherosclerosis; hypertension;
                                                                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                                                                                           Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                               2; Indels
                                                                                                                                                                                                                                                            Sequence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Forward RT-PCR primer for human NOV33b/c/d set 3.
                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                             1495 GGCCTGGACTACTCCTTC 1512
                                                                                                                                                                                                                                                                                                                                                                                 3 GGCCTGGACTGCTTCTTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001US-0318130P.
2001US-0318430P.
2001US-0318765P.
2001US-0322781P.
2001US-0322816P.
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2001US-0323631P.
2001US-0323636P.
2001US-0324969P.
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2001US-0324990P.
2002US-0357303P.
2002US-0360973P.
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2002US-0367753P.
                                                                                                                                                                                                                                                                                          Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative
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2002US-0381664P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACD06434 standard; DNA; 20
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treating pathologies,
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17-SEP-2001;
17-SEP-2001;
19-SEP-2001;
20-SEP-2001;
20-SEP-2001;
25-SEP-2001;
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26-SEP-2001;
15-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD06434;
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1649
 $$$$$$$$$$$$$$$$$$$$$
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2002US-0381672P

17-MAY-2002;

Immunostimulatory nucleic acid #301.

25-SEP-2003 (first entry)

ACD99615;

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The invention relates to an isolated polypeptide comprising one of 127 sequences (appearing as ABO1288-ABO1414) designated as NOVX, a mature form of NOVX, an amino acid sequence which is at least 95% identical to NOVX or an order of Sequence comprising one or more conservative and sequence comprising one or more conservative substitutions in NOVX. Also included are nucleic acids encoding NOVX proteins, determining the presence or amount of Antibody bound to the polypeptide, and determining the presence or amount of antibody bound to the polypeptide, determining the presence of amount of antibody bound to the polypeptide, determining the presence of amount of antibody bound to the polypeptide, determining the presence of cor predisposition to a disease associated with altered levels of cor predisposition to a disease associated with altered levels of cor predisposition to a pathology related to aberrant expression of NOVX is dentifying a potential therapeutic agent to physiological interactions of NOVX, screening for a modulator of activity of or of latency or predisposition to a pathology secociated with NOVX, a vector comprising NOVX DNA, a cell comprising the vector (used to produce NOVX) and an anti-NOVX antibody. The NOVX nucleic acids and polypeptides are useful as a marker for cell or tissue type, and in diagnosing and treating pathologies, diseases, conditions or disorders associated with chromic congenital heart defects, prostate cancer, diabetes, metabolic disorders, neoplasm, graft versus host disease, AlDS, bronchial asthma, Crohn's disease, multiple sclerosis, infectious diseases, anorexia, cancer. associated achexia, neurodegenerative disorders (associated with chromics diseases, Albse may also be used to screen for molecules which inhibit or enhance NOVX activity or function, and detecting sequence is a reverse companied that forense to present experse to the present sequence is a reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptides and polynucleotides having properties related to estimulation of biochemical or physiological responses in a cell or tissue, useful for diagnosing or preventing e.g. atherosclerosis, hypertension, prostate cancer.
                                                                                                                                                                                Zhong M, Li L, Gorman L, Spytek KA, Kekuda R, Taupier RJ; Anderson DW, Vernet CAM, Catleaton E, Miller CE, Shenoy SG; Patturajan M, Pena CEA, Tchernev VT, Padigaru M, Gusev VY; Malyankar UM, Burgess CE, Gerlach VL, Casman SJ, Rieger DK; Grosse WM, Smithson G, Peyman JA, Starling G, Rochenberg ME, Larochelle WJ, Shinmets RA, Crabtree J, Rastelli L, Voss EZ; Boldog FL, Edinger SR, Millet I, Macdougall JR, Ellerman K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example C; Page 473; 849pp; English.
28-WAY-2002; 2002US-0383651P.
29-MAY-2002; 2002US-0384012P.
19-JUN-2002; 2002US-0390155P.
06-SEP-2002; 2002US-00390155.
                                                                                                                                                                                                                                                                                                                                Boldog FL, Edinger SR,
Chapoval A;
                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-313246/30.
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transcriptase (RT)-PCR primer used to assess the tissue specific expression of mRNA encoding a NOVX protein
                                                                                                                                              Sequence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other;
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Gaps
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0.4%; Score 14.8; DB 1; Length 20; 18.9%; Pred. No. 1.7e+03;
                           2; Indels
                           0; Mismatches
                                                      1495 GGCCTGGACTACTCCTTC 1512
            88.9%;
                           16; Conservative
              Best Local Similarity
Query Match
                         Matches
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20

GGCCTGGACTGCTTCTTC

Human; neural crest stem cell line; transplantation; cell therapy; neurological disease; HNC10 cell; neuroprotrotective; cerebroprotective;

PCR; primer; ss.

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WO2003054202-A1 Homo sapiens.

03-JUL-2003

Human HNC10 cell TrkA gene PCR primer #2

15-OCT-2003 (first entry)

ACF04051;

ACF04051/c ID ACF04051 standard; DNA; 20 BP.

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RESULT 1650
ACD99615/c
ID ACD99615 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                             antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                               mmunostimulatory; antiinflammatory; dermatological; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 16; 229pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
                                                                                                                                                                                                                                                                                   29-MAR-2002; 2002US-00112653
                                                                                                                                                                                                                                                                                                                  29-MAR-2001; 2001US-0279642P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1678 GACTICGGGCTGGCCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 GACTTTGGCCTGGCCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-521815/49.
                                                                                                                                                                                                                                                                                                                                                                                                 Krieg AM, Berg DJ;
                                                                                                                                                                                                                                                                                                                                                   (KRIE/) KRIEG A M.
                                                                                                                                                                                                                                                                                                                                                                    BERG/) BERG D J.
                                                                                                                                                                                                                 JS2003050268-A1.
                                                                                                                                                                                                                                                  13-MAR-2003
                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1651
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Gaps

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The invention relates to antisense compounds targeted to nucleic acids encoding protein kinase A regulatory subunit RII alpha. The antisense compounds are useful for medulating the expression of protein kinase A (PKA) regulatory subunit RII alpha and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human protein kinase A regulatory subunit RII alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ds; allergy; asthma; poly-G nucleic acid; aerosol formulation;
hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 9 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fouron Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunostimulatory nucleic acid #299.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 9; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                     1128 GGCCAATGTCTCCGAGCT 1145
                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 GGCCACTGTCTCCGCGCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
                                                                                                                                                                                                                                                                                                              0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-FEB-2000; 2000US-0179991P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB36685 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bratzler RL, Petersen DM,
                                                                                                                                                                                                                       inhibitory oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 88.9
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.4
Best Local Similarity 88.9
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BRAT/) BRATZLER R L. (PETE/) PETERSEN D M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-657977/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2003087848-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB36685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB36685,
*88888888888888
                                                                                                                                                                                                                                                                                                                                                                                                                                               셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ઠે
                                                                                                                                                                                                                                                                                  New primordial human neural crest stem cell having a pluripotent and self -renewing properties, useful for implantation in vivo for cell therapy treatment of human neurological disorders and diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a primordial human neural crest stem cell line suitable for on-demand implantation in vivo into a living host subject comprising a pluripotent and self-renewing neural crest stem cell of human origin. The cell line is useful in the cell therapy treatment of human neurological disorders and diseases. The present sequence is a PCR primer used to isolate human genes from the HNC10 cell line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compounds, useful for modulating the expression of protein
kinase A (PKA) regulatory subunit RII alpha, and for treating a disease
or condition associated with expression of PKA regulatory subunit RII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PKA regulatory subunit RII alpha inhibitory oligonucleotide ISIS102773.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ss; antisense therapy; infection; inflammation; tumour; protein kinase A regulatory subunit RII alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                           Disclosure, Page 39, 70pp, English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1123 ACGCTGGCCAATGTCTCC 1140
                                                                                                    UNIV BRITISH COLUMBIA.
CHILDRENS MEDICAL CENT
UNIV PENNSYLVANIA.
                                                                                                                                                                                                Wolfe JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                  25-APR-2001; 2001WO-US013354.
                                                             05-MAY-2000; 2000US-00565339.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-SEP-2001; 2001US-00954560.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-SEP-2001; 2001US-00954560.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACD44750 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert LM;
                                                                                                                                                                                                Snyder EY,
                                                                                                                                                                                                                                          WPI; 2003-559151/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-511923/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6524854-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACD44750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                            (CHIL-)
(UYPE-)
                                                                                                                                                                                              Kim SU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1652
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Matches

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Gaps

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1678 GACTTCGGGCTGGCCCGG 1695

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Claim 15; Col 43-44; 35pp; English.

20 CTGCCAGGGGAGGAG 3

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The invention relates to a modified oligonuclectide comprising several covalently bound nucleosides including a ribose or decoxyribose sugar portion and a base portion. The nucleosides are joined together by internucleoside linkages such that the base portion of the nucleosides continued in the nucleosides includes a condition of the nucleosides includes a conditionance in maked base sequence. At least one of the nucleosides includes a conditionance of the invention are useful as therapeutics, diagnostics and research agents e.g. for the treatment of various viruses (e.g. AnDS), for modulating the production of proteins by an organism, treating an organism having a disease involving an undesired production of a protein (e.g. atherosclerosis, cancer), detecting the presence or absence of abnormal RNA molecules, or abnormal or inappropriate expression of normal RNA molecules in organisms or cells, and for the selective binding of RNA for use as research reagents and diagnostic agents. The compounds have improved stability to enzymatic degradation with various intracellular and extracellular nucleases, and improved ability to bind to a specific DNA or RNA with fidelity compared to wilder type DNA-DNA and RNA-DNA wild believed and phosphorus-addition of the propertice of the condition of condition of the condition of condition of the condition of conditions of the condition of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modified oligonucleotides useful as therapeutics, diagnostics and research agents comprises several covalently bound nucleosides joined by internucleoside linkages.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            duplexes containing methylphosphonates, phosphoramidates and phosphate triesters. The present sequence is an antisense oligonucleotide of the invention targeting human c-Raf.
                                                                                                                                                                                                                                                                                                                                                                          Human; 88; antisense; c-Raf; virucide; anti-HIV; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; 2'-fluoro substituent; AIDS; atherosclerosis; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match

0.4%; Score 14.8; DB 1; Length 20;
Best Local Similarity 88.9%; Pred. No. 1.7e+03;
Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                        Human c-Raf antisense oligonucleotide #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 31; SEQ ID NO 11; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90US-00463358.
90US-00566977.
92US-00835932.
95US-00468037.
99US-00389283.
20 GACTTTGGCCTGGCCCGG 3
                                                                                                                                                             ADD44694 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JAN-2003; 2003US-00352586
                                                                                                                                                                                                                                                                      15-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cook PD, Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-831271/77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003187240-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-JAN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-AUG-1990;
05-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-OCT-2003,
                                                                                                                                                                                                                  ADD44694;
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The invention relates to a novel method for amplifying a DNA using polymerase chain reaction (PCR) comprising synthesising the first region of a base sequence to be amplified by designing a pair of primers so to place the first region between them and to anneal each other at the 3'-end and carrying out a polymerase chain reaction (PCR) using the primers. Subsequently, the second region is synthesised by designing a pair of primers so as to place the second region partly overlapping with the first region of the base sequence between them and to anneal each other at the 3'-end and carrying out a PCR using the primers. Finally, the first region is annealed to the second region generating the template to carry out a PCR and thus to synthesize a base sequence containing the first and the second regions. The method of the invention may be useful for amplifying a DNA sequence. The current sequence is that of the DNA amplification-related PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Amplification of a DNA, a gene encoding the repeated sequence of an amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HRAS; quadruplex DNA; gene therapy; cancer; cytostatic; oncogene; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                    DNA amplification-related PCR primer - SEQ ID 40.
                                                                                                                                                                                                                                                                                                                                                               (DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU SO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HRAS gene regulatory region quadruplex DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; SEQ ID NO 40; 33pp; Japanese.
                                                                                                                                                     PCR; DNA amplification; 88; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3192 TGCCCCGGAGCTGGAGGA 3209
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                                                                                                                                                                                                                                                                                             29-JUN-2001; 2001JP-00197813.
                                                                                                                                                                                                                                                                                                                              29-JUN-2000; 2000JP-00196242.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match

Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE86164 standard; DNA; 20
          ADD68683 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-375838/36
                                                                                                                                                                                                                        JP2002315583-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        acid sequence.
                                                                                15-JAN-2004
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                                                                                                                                                                                                                                                           29-OCT-2002
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                                               ADD68683;
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ADD68683
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Gaps

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Farrell TJ;

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The present sequence is from the upstream regulatory region of the RET. regulates transcription. The sequence provides an example of intramolecular chair quadruplex Structure ADB86192 with 2 stable tetrads that regulates transcription. The sequence provides an example of intramolecular chair quadruplex DNA structures that have been identified as oncogene regulators. Certain mutations in quadruplex forming mucleotides sequences have been shown to destabilise quadruplex structure and are associated with cancer. Methods are provided for identifying quadruplex mucleotide sequences having destabilising guanine substitutions, for determining whether a subject is at risk of developing or having cancer, pharmacogenomic methods for targeting appropriate prevention or therapoutic regiments, methods for screening molecules that interact with stabilised and destabilised quadruplexes, and therapoutic methods for treating cancers, such as antisense mucleic acid cancer for the appropriate provention or prevention and settabilised and adstabilised quadruplexes, and therapout in the regorifically targets DNA in subjects having a quadruplex-
                                                                                                                                                                                                                                                                                                Identifying a compound that modulates the biological activity of a native quadruplex DNA for treating colorectal cancer comprises determining the presence or absence of interaction between the candidate compound and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; primer; human; PCR; WNT; chronic rheumatoid arthritis; WNT10B; rheumatoid arthritis; osteoarthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indel8
                                                                                                                                                                                                     Siddiqui-Jain A, Grand CL, Bearss DJ, Hurley LH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 3 C; 17 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human WNT2B forward PCR primer SEQ ID NO:5.
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                                                                                                                              CYTERNEX INC. ARIZONA BOARD OF REGENTS.
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                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 3; Page 46; 69pp; English.
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                                05-APR-2002; 2002US-0370358P.
20-AUG-2002; 2002US-0404966P.
20-MAR-2003; 2003US-0456637P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 88.9 es 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        destabilising mutation.
                                                                                                                                                                                                                                                  WPI; 2003-853947/79.
                                                                                                                                                                                                                                                                                                                                                                            test quadruplex DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                              (CYTE-)
                                                                                                                                                          (ARIZ-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1658
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is from the upstream regulatory region of the HRAS gene. The sequence is involved in the regulation of transcription. It forms a quadruplex structure through the formation of guanine tetrads. The sequence provides an example of intramplecular chair quadruplex DNA structures that have been identified as oncogene regulators. Certain mutactions in quadruplex forming nucleotides sequences have been shown to destabilising quadruplex structure and are associated with cancer. Methods are provided for identifying quadruplex nucleotide sequences having destabilising guanine substitutions, for determining whether a subject is at risk of developing or having cancer, pharmacogenomic methods for targeting appropriate prevention or therapeutic regiments, methods for screening molecules that interact with stabilised and destabilised quadruplexes, and therapeutic methods for treating cancers, such as antisense nucleic acid cancer therapy that specifically targets DNA in subjects having a quadruplex-destabilising mutation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying a compound that modulates the biological activity of a native quadruplex DNA for treating colorectal cancer comprises determining the presence or absence of interaction between the candidate compound and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Platelet derived growth factor alpha; PDGF; quadruplex DNA; gene therapy; cancer; cytostatic; oncogene; human; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                          Siddiqui-Jain A, Grand CL, Bearss DJ, Hurley LH, Farrell TJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 3 C; 17 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RET gene regulatory region quadruplex DNA.
                                                                                                                                                                                                                                                                                   (CYTE-) CYTERNEX INC.
(ARIZ-) ARIZONA BOARD OF REGENTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 46; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2920 GGGCGGGCGTGGGGGG 2937
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20-MAR-2003; 2003US-0456637P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE86160 standard; DNA; 20 BP.
                                                                                                                                                                                           05-APR-2002; 2002US-0370358P.
                                                                                                                                           04-APR-2003; 2003WO-US010658
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 88.95
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-853947/79.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    test quadruplex DNA.
                                             WO2003087317-A2
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Homo sapiens.
                                                                                             23-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumors, using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a method of preventing or treating tumour metastasis in an animal comprising administering to the animal an oligonucleotide 8-50 nucleotides in length, which is targeted to mRNA encoding human raf and capable of inhibiting raf expression. Also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour metastasis; human; raf; raf expression inhibitor; cytostatic; antiarteriosclerotic; antisense-therapy; hyperproliferative disorder; atherosclerosis; tumour; c-raf kinase; antisense oligonucleotide; ss.
                                                        Detection of over expression of WNT10B by analysis of synovial fluid,
joint tissue or peripheral blood for diagnosis of chronic rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
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such as hyperproliferative disorders, atherosclerosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human c-raf kinase antisense oligonucleotide seq id 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 3 C; 8 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                             Disclosure; SEQ ID NO 5; 28pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTGAGTCTGGGCAGCTGT 3648
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96US-00756806.
97US-00888982.
98WO-US013961.
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                                                                                                                                                                                                                                                                                                                                                                                             primer used in the invention.
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2000US-00506073.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
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nes 16; Conservative
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WPI; 2003-854488/79
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18-FEB-2000;
25-JAN-2002;
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26-NOV-1996;
07-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUN-2002;
                                                           Detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-FEB-2004
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ADF09
ADF0
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disclosed are raf oligonucleotides, nucleic acids, proteins and compositions used in the methods of the invention. The oligonucleotides have cytostatic and antiatrateriosclerotic properties, are useful as Rafinhibitors and in antisense-therapy. The methods and compositions of the present invention are useful for preventing and/or treating conditions associated with raf expression, such as hyperproliferative disorders, atherosclerosis and tumours. This sequence represents a human c-raf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nouse; murine; PCR; ss; vanilloid receptor; VR; pain perception; TRPV3; VRLS; VRLX; VR4; TRPV7; TRPV4; VRL3; OTRPC4; TRPM8; TRPX; trkA+; inflammation; skin disorder; cancer; analgesic; antiinflammatory;
                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bevan S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiinflammatory, dermatological and cytostatic activities. This
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                                                                                                                                                                                       Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
                                                                                                                                                                                                                           Indels
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                                                                                                                                                     Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peier A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine TRPV transcript PCR primer SegID 48.
                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            dermatological; cytostatic; primer.
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                                                                                                                     kinase antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , 2001US-0297835P.
2002US-0351238P.
2002US-0352914P.
2002US-0357161P.
2002US-0381086P.
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                                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                         0.4%;
                                                                                                                                                                                                                                                         844 CTGCCAGCCGAGGAGGAG
                                                                                                                                                                                                                                                                                             20 CTGCCAGGGGAGGAGGAG
                                                                                                                                                                                                                                                                                                                                                                                 ADG32593 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                     Query Match
Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-JUN-2001; 22-JAN-2002; 29-JAN-2002; 212-FEB-2002; 215-MAY-2002; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                    ADG32593;
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                                                                                                                                                                                                                                                                                                                                                 RESULT 1660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mouse;
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0.4%; Score 14.8; DB 1; Length 20;
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Query Match

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acids and encoded proteins thereof. Specifically, it refers to certain members of the WR family that are involved in pain perception, in particular, TRPU3 (previously known as VRLS, VRLX, VR4 & TRPV3), TRPV4 (previously known as VRL) & OTRPC4) and TRPW3 (previously known as TRPX). Furthermore, this invention includes trKA+ pain specific genes expressed in the sensory neurons of the dorsal root ganglia. Accordingly, such compositions can be useful for the diagnosis, treatment and prevention of pain, inflammation, skin disorders and cancer, and so exhibit analgesic, antiinflammatory, dermatological and cytostatic activities. This colionucleotide sequence is a PCR primer used to amplify the murine TRPV3
oligonucleotide sequence is a PCR primer used to amplify the murine TRPV3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mouse; murine; PCR; ss; vanilloid receptor; VR; pain perception; TRPV3; VRLS; VRLX; VR4; TRPV7; TRPV4; VRL3; OTRPC4; TRPM8; TRPX; trkA+; inflammation; skin disorder; cancer; analgesic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to novel vanilloid receptor (VR) related nucleic
                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated TRPV3, TRPV4 or TRPW8 vanilloid receptor nucleic acid molecule and polypeptides, useful for the diagnosis and treatment of disorders such as pain, inflammation, skin diseases and cancer.
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                                                                                                              Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcintyre P,
                                                                   Sequence 20 BP; 7 A; 1 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 20 BP; 2 A; 10 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peier A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Murine TRPV transcript PCR primer SeqID 73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 73; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dermatological; cytostatic; primer.
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                                                                                                                                                                                                                  186 GGAGGACGAGGCTGAGGA 203
                                                                                                                                                                                                                                                                 GGAGGACGAAGGTGAGGA 19
                                                                                                                                                                                                                                                                                                                                                                                    ADG32618 standard; DNA; 20 BP
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12-FEB-2002; 2002US-035746IP.
15-MAX-2002; 2002US-0381086P.
16-MAX-2002; 2002US-0381739P.
                                                                                                                   Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
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2002US-0351238P
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                         DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-156962/15.
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22-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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SXXS
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The invention relates to a novel polymucleotide isolated and purified from a human gene having any one of 935 fully defined sequences as given in specification, or a sequence having a base substitution. The invention curther relates to: an oligomucleotide containing single nucleotide polymorphisms; a PCR primer set chosen from the combination of two DNA fragments from any one of 1220 fully defined sequences as given in comparing the from any one of 1220 fully defined sequences as given in a microarray equipped with the SNP containing oligo. The isolated human contained invention is useful for detecting the single nucleotide plymorphisms in human gene. The isolated human gene is also useful for diagnosis of disease and determination of side effect to a medical agent. The isolated human gene is also effective in detecting single nucleotide polymorphism in a human gene. This polynucleotide sequence represents one of the PCR primers used in the single nucleotide polymorphism cone of the invention.
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                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human; single nucleotide polymorphism; microarray; side effect; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism detection primer, SEQ ID No 1612.
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                                                           Indels
88.9%; Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; SEQ ID NO 1612; 704pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1292 CCGTGAAGATGCTGAAAG 1309
                                                                                                              203
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                                                                                                                                                   19 GGAGGACGAAGGTGAGGA 2
                                                                                                                                                                                                                                                                                                         BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-FEB-2002; 2002JP-00034717.
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                                                                                                              186 GGAGGACGAGGCTGAGGA
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                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
                                                        16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                               ADF88029;
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                                                                                                                                                                                                                                                   RESULT 1662
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ID ADI120
XX
AC ADI120
                                                           Matches
                                                                                                                                                                                                                                                                                  ADF88029
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antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antianthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Human; antisense; lung dysfunction; nasal airway dysfunction;

Human oligonucleotide sequence.

(first entry)

17-OCT-2003

ABZ90613;

ВЪ.

ABZ90613 standard; DNA; 20

ABZ90613/c

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ss; nuclease resistant; mixed sequence; 2'-deoxyfuranosyl; c-raf;
                                                                                                                  /mod_base= OTHER
/note= "OTHER= Optionally 2'-O-methyl"
                   Human c-raf antisense oligonucleotide ISIS #7854
                                                                                                /note= "OTHER= Phosphorothioate"
                                                                                                                                                                                                                                                                                           Example 31; SEQ ID NO 11; 48pp; English.
                                                                      Location/Qualifiers
                                                                                         base= OTHER
                                                                                                                                                                                    92US-00835932.
92US-00835932.
                                                                                                                                                                 99US-00389283
                                                                                                                                                                              90US-00463358
                                                                                                                                                                                                       95US-00468037
98US-00035357
                                                                                   tag= b
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      (first entry)
                                                                                                             /*tag=
                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                        Cook PD, Kawasaki AM;
                                                                                           pom/
                                                                                                                                                                                                                                                     WPI; 2003-566474/53.
                                      antisense; human
                                                                      Key
modified_base
                                                                                                      modified base
                                                                                                                                       US6531584-B1.
                                                   Homo sapiens
                                                                                                                                                                                    13-AUG-1990;
05-MAR-1992;
01-JUL-1992;
                                                                                                                                                                 02-SEP-1999;
      15-APR-2004
                                                                                                                                                    11-MAR-2003.
                                                                                                                                                                                                               35-MAR-1998;
                                                                                                                                                                              11-JAN-1990
                                                                                                                                                                                                        06-JUN-1995
                                                         Synthetic
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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

Disclosure; SEQ ID NO 5855; 872pp; English.

ubiquinone.

Katz E, Pabalan J, Aguilar D;

Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S;

Miller S,

Nyce JW.

WPI; 2003-229219/22.

23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC

WO200285308-A2. sapiens.

Ношо

31-OCT-2002.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation condon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubjudinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of denosine of increasing levels of denosine of increasing levels of denosine or levels of denosine or 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lung surfactant in a subject's tissue, or treating bronchoconstriction, uning inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequences
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specification, but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
The invention relates to a nuclease resistant mixture organization collections comprising at least one modified 2'-deoxyfuranosyl oligonucleotides comprising at least one modified 2'-deoxyfuranosyl group. The modified oligonucleotides are disclosed as being useful for modulating the production of a protein by an organism, and especially for treating a disease in an organism which is characterised by the undesired production of a protein. The oligonucleotides may be used to treat diseases caused by viruses or other agents. The oligonucleotides may also be used for diagnostic methods for detecting the presence or absence of abnormal RNA molecules in an organism or cell. Oligonucleotides of the normal RNA molecules in an organism or cell. Oligonucleotides of the invention that selectively bind RNA may also be useful as research reagents. The new oligonucleotides are nuclease resistant and hybridise to RNA or DNA targets with high strength and specificity. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nuclease resistant mixed sequence oligonucleotides useful as therapeutics, diagnostics, and research agents comprise at least one modified 2'-deoxyfuranosyl group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to RNA or DNA targets with high strength and specificity. The sequence represents a human c-raf antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
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Gaps
                                                                            .;
0
                                0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels
Sequence 20 BP; 1 A; 1 C; 13 G; 5 T; 0 U; 0 Other;
                                                                                                                2159 CCCCGGCCCCACCAGCA 2176
                                                                                                                                         18 CCCCACCCACCCAGCA 1
                                                      Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                          6.
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RESULT 1665

844 CTGCCAGCCGAGGAG 861

CTGCCAGGGGAGGAGGAG

20

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RESULT 1664

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubjudinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an exceptor, producing bronchedilation, increasing levels of adenosine receptor, producing bronchedilation, increasing levels of adenosine receptor, producing bronchedilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating broncheconstriction, lung inflammation, lung allergies, or a respiratory disease or condition, note: The sequence data for this patent is not represented in the printed especialisation, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                            Human, antisense, lung dysfunction; nasal airway dysfunction; antianflammatory steroid; ubiquinone; antianflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Katz E, Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 13904; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                      Human tryptase a oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                  ABZ98662 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENESIS PHARM INC.
                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                             WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                          17-0CT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                         31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ubiquinone,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Miller S,
                                                             ABZ98662;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nyce JW,
ABZ98662/c
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                                                                   Gaps
                                                                   ;
0
                                0.4%; Score 14.8; DB 1; Length 20;
38.9%; Pred. No. 1.7e+03;
ive 0; Mismatches 2; Indels
Sequence 20 BP; 2 A; 10 C; 5 G; 3 T; 0 U; 0 Other;
                                                88.9%;
                                                                   16; Conservative
                                                Best Local Similarity
Matches 16; Conserv
                                 Query Match
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18

CGGGAGGCCCGGGGCATG

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3196 CCGGAGCTGGAGGATCCC 3213
                            19 CTGGAGCTGGAGGAGCCC
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomuclectide antisense to the initiation codon, coding region, 5 or 3' end genomic flanking regions, 5 and 3' intron-exon junctions, or regions within 2-10 muclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, invention and the invention and the invention and the invention and the composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of, or reducing sensitivity to adenosine, reducing levels of adenosine, receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed because of the patent is not represented in the printed or the patent is not represented in the printed or the patent is not represented in the printed or the patent is not represented in the printed or the patent is not represented in the printed or the patent is not the printed in the printed or the patent is not trepresented in the printed or the patent is not trepresented in the printed or the patent is not trepresented in the printed or the patent is not trepresented in the printed or the patent is not trepresented in the printed or the patent is not the presented in the printed or the patent is not trepresented in the printed or the patent is not the patent in the printed or the patent in the printed or the patent is not the patent in the printed or the patent in the printed or the patent in the printed or the patent in the patent in the patent in the patent in the patent 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                    antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D;
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                             Human; antisense; lung dysfunction; nasal airway dysfunction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 6 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 14092; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specification, but was obtained in electron: at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                 lung inflammation; respiratory disease; ds.
                                                                                                                                                                                       Human PDE4A oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1573 CAGGTGGCCCGGGGCATG 1590
                      ABZ98850 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%;
                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nyce JW, 1
                                                                             ABZ98850;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Loca
Matches
4BZ98850
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RESULT 1667

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction as pecond active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of composition in receptor, producing bronchodilation, increasing levels of adenosine or receptor, producing bronchodilation, increasing levels of biquinone or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                surfactant in a subject's tissue, or treating bronchoconstriction, inflammation, lung allergies, or a respiratory disease or condition: The sequence data for this patent is not represented in the printed iffication, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                               antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiantasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Katz E, Pabalan J, Aguilar D;
                                                                                                                                                                                           antisense; lung dysfunction; nasal airway dysfunction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure, SEQ ID NO 3439; 872pp; English.
                                                                                                                                                                                                                                                                                                   lung inflammation; respiratory disease; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                   Human oligonucleotide sequence.
                      ABZ88197 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPIG-) EPIGENESIS PHARM INC
                                                                                                 17-OCT-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              specification, but
                                                                                                                                                                                                                                                                                                                                                                                       WO200285308-A2
                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                              31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Miller S,
                                                                  ABZ88197;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nyce JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lung
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Gaps ö 0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indel8 88.9%; 16; Conservative Best Local Similarity Query Match Matches

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RESULT 1668

Human; antisense; lung dysfunction; nasal airway dysfunction; Human oligonucleotide sequence. ВР. ABZ87781 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ87781; ABZ87781/c

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

sapiens. Ношо WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, Sandrasagra A, K i, Shahabuddin S; Li Y, Sar Tang L, Willer S, 3

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 3023; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 and genomic flanking regions, 5' and genomic flanking regions, 5' and genomic flanking regions, 6' and all intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or mailgnant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing shorthaltation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung suffammation, lung allergies, or a respiratory disease or condition Note: The sequence data for this spatent is not represented in the printe specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 1 A; 9 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ; 0 0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels Pred. No. 1.76 0; Mismatches 2114 CCAGCTCCTCAGGGACG 2131 88.98; Local Similarity 88.5 Query Match Best Loca Matches ∴∂

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19 CCAGCTGCCCAGGGGACG

RESULT 1669

Gaps

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example from the present invention
                                                                                                                                                                                                        Claim 3; Page 89; 167pp; English.
                                                                                                                                            15-JUL-2002; 2002WO-US022672.
                                                                                                                                                    26-JUL-2001; 2001US-00915814.
   ABZ82743 standard; DNA; 20
                   (first entry)
                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                     Watt AT,
                                                                                                                                                                            WPI; 2003-239411/23.
                                                                                                                            WO2003010139-A2
                                                                    Key
modified_base
                                                                                       modified_base
                                                                                                        modified base
                                                        Homo sapiens
Synthetic.
                   14-MAY-2003
                                                                                                                                    06-FEB-2003
                                                                                                                                                                    Butler MM,
            ABZ82743;
ABZ82743/c
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Query Match Best Local Similarity

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*tag=

/mod

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16. .20 /*tag=

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The invention relates to antisense compounds targetted to a nucleic acid encoding thyroid hormone receptor interactor 6 (TRIP6) to inhibit its expression. TRIP6 is also known as OPA-interacting protein-1 (OIP-1) and zyxin-related protein-1 (OIP-1) and zyxin-related protein-1 (OIP-1) and hitisense compounds of the invention are useful for modulating the expression of TRIP6 and for treating diseases or conditions associated with the expression of TRIP6 such as hyperproliferative disporders (e.g. cancer). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as tresearch reagents and kits and in distinguishing between functions of various members of a biological pathway. The are also useful in antisense therapy. The present sequence is an antisense oligo targetted to human TRIP6 DNA. This oligo is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides targeted to nucleic acids encoding thyroid hormone receptor interactor 6, useful for diagnosing or treating hyperproliferative disorders, such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
1. .20
/*tag= a
/*tag= a
/*mod_base= OTHER
/noce= "Phosphorothioate backbone; All cytidine residues
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense; thyroid hormone receptor interactor 6; TRIF6; tumour; OPA-interacting protein-1; OIP-1; zyxin-related protein-1; prophylaxis; inflammation; therapy; hyperproliferative disorder; infection; cancer; chromosome 7q22; ZRP-1; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
Indels
                                                                                                                                                                                                                                                                                                                                                   Human TRIP6 antisense oligonucleotide ISIS #198770.
5
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; Page 76; 111pp; English.
                                           2705 CCCTGCCCTCAGAGACT 2722
                                                                                       7
                                                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-NOV-2001; 2001US-00008789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-NOV-2002; 2002WO-US035479.
                                                                                  19 CCTGCTCCTCCAGACT
                                                                                                                                                                                         898/c
ACC82898 standard; DNA; 20
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/*tag≈ b
                                                                                                                                                                                                                                                                                                       (first entry)
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16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dobie K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-430662/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003040328-AZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
                                                                                                                                                                                                                                                                                                    27-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                        ACC82898;
                                                                                                                                                              RESULT 1670
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                  ACC82898,
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                                                                                     셤
                                                                                                                                                                                                                                  The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding a hormone-sensitive lipase (HSL) or a splice variant of HSL. The compound specifically hybridises with and inhibits the expression of HSL or a splice variant of HSL, or specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding HSL. (1) have anorectic, antidiabetic and cytostatic activities, and can be used in antisense therapy. (1) is useful for tracting an animal, particularly human, suspected of having an abnormal metabolic condition such is obesity,
                                                                                                                                                                                                Hormone-sensitive lipase; antisense oligonucleotide; inhibitor; obesity; phosphorothioate; antidiabetic; anorectic; cytostatic; antisense therapy; abnormal metabolic condition; hyperlipidaemia; type 2 diabetes; cancer; hyperproliferative disorder; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hyperlipidaemia, type 2 diabetes, a hyperproliferative disorder such as cancer (e.g. pituitary, colorectal, breast, testicular, pulmonary or epithelial cancer). (I) is also useful in modulating blood glucose levels, particularly plasma or serum glucose levels, in a diabetic animal. The present sequence represents a human hormone-sensitive lipase chimeric phosphorothicate antisense oligonucleotide, which is used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides targeted against nucleic acids encoding hormone-sensitive lipase, useful for treating abnormal metabolic condition, e.g. hyperlipidemia and obesity, or a hyperproliferative disorder, e.g. cancer.
                                                                                                                                                            Human HSL chimeric phosphorothioate oligonucleotide SEQ ID NO:132.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOE) wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOE) wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'note = "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 5 A; 4 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1;
88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
1. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 base= OTHER
                     ВЪ
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Gaps

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Indela

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various cell functions and physiological processes and conditions and for
             diagnosing conditions associated with raf expression and for research purposes. The present sequence is an antisense oligonucleotide targeting a human raf mRNA
                                                                                                 0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03;
                                                                     Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             that may be useful as herbicides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (DUPO ) DU PONT DE NEMOURS & CO (PRAB/) PRABHU V.
                                                                                                                                                           861
                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-JUN-2001; 2001US-0297198P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JUN-2002; 2002WO-US018230.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%;
                                                                                                                                                           844 CTGCCAGCCGAGGAG
                                                                                                                                                                                       20 CTGCCAGGGGAGGAG
                                                                                                                88.98;
                                                                                                                                                                                                                                                             AADS5037 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                               Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Damude HG, Flint D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-201331/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2002101013-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Intermediates.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Alstroemeria.
                                                                                                                                                                                                                                                                                                                      26-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-DEC-2002.
                                                                                                                                                                                                                                                                                            AAD55037;
                                                                                                  Query Match
                                                                                                                                                                                                                                  RESULT 1672
                                                                                                                                                                                                                                                AAD55037
                                                                                                                                                                                                                                                                             888888
                                                                                                                                                            8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an oligonucleotide 8-50 nucleotides in length which is targeted to mRNA encoding human c-raf, a-raf or b-raf (raf is a protein kinase playing a regulatory role in signal transduction, regulating cell proliferation and has been implicated in lung carcinoma), and which is capable of inhibiting raf expression. Also included is a composition comprising the oligonucleotide and a pharmaceutically acceptable carrier. The antisense oligonucleotide is useful for inhibiting the expression of human raf in human cells or tissues, by contacting the human cells or tissues with the oligo. The oligo. Is also is useful for treating or preventing a disease or condition associated with the expression of raf by administering it in combination with a chemcherapeutic agent to a human or cells of the human, where the expression of raf is abnormal expression, and the condition is a hyperproliferative condition such as cancer, anglogenesis or neovascularisation (preferably ocular anglogenesis or neovascularisation). The oligo. is also useful for inhibiting hyperproliferation of cells. The oligos are also useful as tools, for example for detecting and determining the role of raf expression in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense oligonucleotide which is targeted to mRNA encoding human
                                                                                                                                                                                                                                                                                                                    Human; ss; antisense; c-raf; a-raf; b-raf; protein kinase; cancer; signal transduction; cell proliferation; lung carcinoma; cytostatic; antisense gene therapy; chemotherapeutic agent; angiogenesis; hyperproliferative condition; neovascularisation; ocular anglogenesis.
                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      raf and which is capable of inhibiting raf expression, useful for treating or preventing hyperproliferative conditions such as cancer.
                                                                     ö
                                         0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                         Antisense oligonucleotide targeting human c-raf, ISIS7854.
             G; S T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 8; 42pp; English.
                                                                                                  1116 GGCCCCCACGCTGGCCAA 1133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95WO-US007111.
96US-00756806.
97US-00888982.
98WO-US013961.
98US-00143214.
                                                                                                                                                                                                     ACD42098 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JAN-2002; 2002US-00057550
             Sequence 20 BP; 2 A; 5 C; 8
                                                                                                                     18 GAGCACCACGCTGGCCAA
                                                                                                                                                                                                                                                             (first entry)
                                                                     16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-503332/47.
                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MONI/) MONIA B P.
                                                                                                                                                                                                                                                                                                                                                                                                                          US2003032607-A1.
                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-MAY-1995;
26-NOV-1996;
07-JUL-1997;
                                                                                                                                                                                                                                                             05-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-AUG-1998;
18-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           36-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monia BP;
                                                                                                                                                                                                                                  ACD42098;
                                          Query Match
                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated nucleic acid fragment encoding a tuliposide A synthesizing protein, useful for creating recombinant organisms that have the ability to synthesize tulipalin A, tuliposide A or tuliposide A pathway
                                                                              Alpha-methylene-gamma-butyrolactone; glutamate decarboxylase; herbicide; enzyme; gamma-aminobutyrate aminotransferase; UDP-glucosyltransferase; gamma-hydroxybutyrate dehydrogenase; tulipalin A; plant; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence is a primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to clone Alstroemeria glutamate decarboxylase homologue gene (gad3)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present
Alstroemeria gad3 gene cloning primer, NW12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prabhu V, Wang H;
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1.
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885 CAGTGTGTATGCAGGCAT 902
  CATTGTGTATGCAGGAAT 18
                ABD24011 standard; DNA; 20
                                                                                                        bronchodilating agent.
                                                                                            WPI; 2003-093058/08.
                                                             WO200285309-A2
                          29-JUL-2004
                                                                  31-OCT-2002.
                                                                                        Miller S,
                     ABD24011;
                                                                                     Nyce JW,
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating construction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cuffactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production congrises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoconstriction, allergies asthma, impeded respiration, respiratory cinflammation, allergies asthma, impeded respiration, respiratory clintammation, allergies asthma, impeded respiration, respiratory clintlammation, allergies asthma, impeded respiration, respiratory clintammation, emphysema, chronic observations pulmonary disease, pulmonary infections, bronchitis or cancer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; antalgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. t Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted uncleic acids associated with lung airway or lung dysfunction, and Aguilar D; Human calmodulin 2-derived oligonucleotide SEQ ID 3023 Katz E, Pabalan J, Claim 15; SEQ ID NO 3023; 763pp; English. Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; ВР 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC (first entry)

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The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide containing less percentage of adenosine, targeted t
nucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                              Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pabalan J,
                                                                                                                        Sequence 20 BP; 1 A; 9 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PDE4A-derived oligonucleotide SEQ ID 14092.
                                                                                                                                                                                                     0; Mismatches
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Tang L, Shahabuddin S;
                                                                                                                                                                                                                                             2114 CCAGCTCCTCAGGGGACG 2131
                                                                                                                                                                                                                                                                            19 CCAGCTGCCCAGGGACG 2
                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                            0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                      ABD31881 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                     Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-093058/08.
                                                                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                            ABD31881;
                                                                                                                                                                Query Match
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B
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                                                                                                                                                                                                                                                                                                                                               RESULT 1
ABD31881
    88688888
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vivlemore401-10.rng

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treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory inflammation, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system or and a surfaces and anti-sense oligonucleotides into products that free adenosine into the system or and a surfaces and anti-sense oligonucleotides into products that free adenosine into the system or and a surfaces and anti-sense oligonucleotides into products that free adenosine into the system or and a surfaces and anti-sense oligonucleotides into products that free adenosine into the system or and a surfaces and anti-sense oligonucleotides and a surfaces and a surfa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unwanted effects due to it
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Sequence 20 BP; 2 A; 6 C; 10 G; 2 T; 0 U; 0 Other;

Gaps .; 0 Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03; 0; Mismatches 2; Indels 88.9%; Matches 16, Conservative Query Match Best Local Similarity

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ઠ 셤 ABD24427 standard; DNA; 20

29-JUL-2004 (first entry) ABD24427;

AI652901-derived oligonucleotide SEQ ID 3439.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. RESULT 1675
ABD24427
XX
AC ABD24427
XX
AC ABD24427
XX
DT 29-JUL-2
XX
C AL652901
XX
C A

Homo sapiens.

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D;

Katz E,

Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; WPI; 2003-093058/08.

Miller S,

Nyce JW,

oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and Pharmaceutical composition for treating asthma, has antisense bronchodilating agent.

Claim 15; SEQ ID NO 3439; 763pp; English.

This invention describes a novel composition (a) a first active agent,

Katz E, Pabalan J, Aguilar D;

Li Y, Sandrasagra A,

Nyce JW,

23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. EPIG-) EPIGENESIS PHARM INC.

31-OCT-2002.

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comprising oligonucleotides, effective for alleviating
bronchoconstriction, respiratory tract inflammation, allergies and
reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
surfactant depletion or hyposecretion, when administered to a mammal. The
oligonucleotides are derived from a gene encoding or regulating
expression of a target polypeptide associated with lung airway or lung
dysfunction or cancer and can be anti-sense to the corresponding mRNA.
The invention also describes a kit, that comprises: (a) a delivery
device, in separate containers, (b) the oligonucleotides, (c) the invention has antiallersy, antialmentory, antiasthmatic,
of the invention has antiallersy, and for use of the kit. The composition
of the invention has antiallersy, and containers of
the invention has antiallersy, and containers of
the invention has antiallersy and cytostatic activity, is a
beta-adrenergic agonist. The composition is useful for preventing or
treating a respiratory, lung or malignant disease. The administered
composition comprises oligo and is administered to reduce the production
or availability, or to increase the degradation of the target mRNA or to
reduce the amount of target polypeptide present in the lungs. The
pulmonary obstruction, and/or bronchoconstriction and/or lung
or inflammation, allergies and/or bronchoconstriction and/or lung
inflammation, allergies as thma, impeded respiration, respiratory
distress syndrome, pain, cystic fibrosis, allergic rhinits, pulmonary
transplantation rejection, pulmonary infections, bronchtis or cancer.
The reduced adenosine content of the anti-sense oligos corresponding to
thymidines present in the target RNA serves to prevent the breakdown of
the oligonucleotides into products that free adenosine into the system
e.g., lung, brain, heart, kidney, ecc, tissue environment and thereby, to
the oligonucleotides into product of it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenargic agonist; respiratory disease; pulmonary vasocometriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Tryptase a-derived oligonucleotide SEQ ID 13904.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              623 CCCACATCCAGTGGCTCA 640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 cccacarcaagaggcrca 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 88.9°
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200285309-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABD31693,
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This invention describes a novel composition (a) a first active agent, comporising oligonuclectides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition or availability, or to increase the degradation of the target mRNA or to increase the administered to reduce the amount of target polypeptide present in the lungs. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstruction, inflammation, allergies, asthem, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emplysem, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                               Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                         Claim 15; SEQ ID NO 13904; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prevent any unwanted effects due to it
Tang L, Shahabuddin S;
                                                                                                                                                                                                   bronchodilating agent.
                                                      WPI; 2003-093058/08
Miller S,
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Sequence 20 BP; 2 A; 10 C; 5 G; 3 T; 0 U; 0 Other;

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Gaps
                                .;
0
Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                              0; Mismatches
                Best Local Similarity 88.9%;
                                  16; Conservative
   Query Match
                                  Matches
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ABD26843 standard; DNA; 20 BP. ABD26843; RESULT 1677

29-JUL-2004 (first entry)

AA278764-derived oligonucleotide SEQ ID 5855.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. ABD26843/ ID ABD2 AC ABD2 AC ABD2 XX ABD2 XX ABD2 XX ABD2 XX Huma XX Huma XX Huma XW resp XW resp XW anal XW resp XW emph XW emph XX EMPH XX ABD2 XM A

Homo sapiens.

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This invention describes a novel composition in a like active agent, comprising oligonucleotides, effective for alleviating critication, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, contractant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating contraction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antialergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a contraction a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or surfactatin the prograduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies and/or surfactatin the present in the lungs. C distress syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary contransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligons corresponding to the continue present in the target RNA serves to prevent the breakdown of the charged adenosine content of the anti-sense oligons or cancer. The reduced adenosine content of the anti-sense oligons or cancer. The reduced adenosine content of the anti-sense oligons or cancer. The charge and content of the anti-sense oligons or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the oligonucleotides into products that free adenosine into the system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                         to
                                                                                                                                                                                                                                                                                                                                                                                                                    oligonuclectide containing less percentage of adenosine, targeted t
nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                       Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 1 C; 13 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                       Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 15; SEQ ID NO 5855; 763pp; English.
                                                                                                                                                                                                                                                                     Sandrasagra A, Ka
L, Shahabuddin S;
                                                                                                                        23-APR-2002; 2002WO-US013143
                                                                                                                                                                      24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                    (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bronchodilating agent
                                                                                                                                                                                                                                                                                                                                            WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                            Tang L,
                                                                                                                                                                                                                                                                     Li Y,
                       WO200285309-A2
                                                                       31-OCT-2002
                                                                                                                                                                                                                                                                                            Willer S,
                                                                                                                                                                                                                                                                       Nyce JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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ö Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03; 0; Mismatches 2; Indels 88.98; 16; Conservative Local Similarity Best Loca Matches

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Gaps

2159 CCCCGCCCCACCCAGCA 2176 18 CCCCACCCCACCCAGCA

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ADH71020 standard; DNA; 20 ADH71020/c ID ADH7100 XX AC ADH7100 XX DT 25-MAR XX

RESULT 1678

ADH71020;

25-MAR-2004 (first entry)

vivlemore401-10.rng

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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers pecifically priming and allowing amplification of each Whete agene.

To specifically priming and allowing amplification of each Whete agene.

To conclude the contract of seases and seases and seases.

To including autoinmune diseases, degenerative nervous system diseases.

To arrophic gastriitis. Degenerative nervous system diseases include multiple actoring and Alzheimer's disease. Hypersensitivity diseases include multiple sclenois and Alzheimer's disease. Hypersensitivities of a sonate. Hypersensitivities such as those present in Coodpasture's syndrome and Type IV hypersensitivities such as those caused by viruses such as HIV. fungal infections actored by viruses such as HOSP caused by cheevest infections caused by the yeast gaung Candida, parasitic infections such as those caused by shistosomes, filaria and bacterial infections such as those caused by shistosomes, filaria and bacterial infections such as those caused by shistosomes, filaria and bacterial infections such as those caused by contracterium. Neoplastic diseases include lymphoproliferative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                           degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; degenerative nervous system disease; multiple sclerosis; allergy; type II hypersensitivity disease; type I hypersensitivity; loodpasture's syndrome; type IV hypersensitivity; looppasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; fillaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer; breast cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a cosmid C215 repeat region PCR
                                            human; T-cell associated disease; Wheta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 7 A; 4 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; SEQ ID NO 1214; 164pp; English.
Cosmid C215 repeat region PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-00263959.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-00309335.
95US-00531241.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-059052/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hood LE, Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HOOD/) HOOD L E. (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                       US2002150891-A1.
                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                    17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vbeta gene.
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The present invention relates to antisense compounds targeted to a nucleic acid encoding G protein-coupled receptor (GPCR) kinase 6 (GRK6). The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of GRK6. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-methoxyethyl (2'-MOE) sugar moiety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methylcytosine. The antisense oligonucleotides are useful for the treatment of diseases such as rheumatoid arthritis, drug addiction, uterine contractility, hypertension, and diseases or conditions arising from abbrrant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding G protein-coupled receptor (GPCR) kinase 6, useful
for treating diabetes, drug addiction, uterine contractility and
                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/mode= "This oligonuclectide has a phosphorothicate
/mote= "This oligonuclectide has a phosphorothicate
backbone and 2'-methyoxyethyl (2'-MOE) wings at the 5'
and 3' ends, which are 5 nuclectides in length at each
end. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaрв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                     Antisense therapy, human, G protein-coupled receptor kinase 6; GPCR kinase 6; GRK6; rheumatoid arthritis; drug addiction; uterine contractility, hypertension; aberrant haematopoiesis; antiinflammatory; antiarthritic; antiirheumatic; hypotensive; phosphorothioate; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             haematopoiesis. The present sequence represents an antisense oligonucleotide used in the examples of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                            Human GRK6 DNA, antisense oligonucleotide #64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; SEQ ID NO 74; 58pp; English.
                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3670 ATGGCTCAGGGTGGTCTC 3687
BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAY-2002; 2002US-00159856.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2002; 2002US-00159856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Argecraagegragrance
ADH48272 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                    1. .20
/*tag= a
                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-052027/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003228689-A1
                                                                                                                                                                                                                                                                                                   Key
modified_base
                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypertension.
                                                                       25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Freier SM,
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                                    ADH48272;
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Gaps

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0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels

3377 TIGCIGIGIGICCCAGGC 3394

Conservative

16;

Matches

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Local Similarity

TIGCICIGICICCCAGGC 1

18

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phosphorothioate backbone; 2'-methoxyethy1; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Crosby SD, Nalseth AE;
                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2004
                                                                    34-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1682
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to nucleic acids encoding a mammalian glucocorricoid receptor. The antiense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNR sequence represents an antisense oligonucleotide that targets the numan glucocorricoid receptor gene. NOTE: The present Boardence contains 2'-methoxyethyl (2'-NOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss;
                                                                                                                                      antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetees; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2-methoxyethyl; 2'-MOB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                  Human glucocorticoid receptor-specific antisense oligonucleotide #2069.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human glucocorticoid receptor-specific antisense oligonucleotide #2426.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 2069; 985pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1928 ACTGCACACACGACCTGT 1945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
                                              ADH65235 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                       20-MAY-2003; 2003WO-US016084.
                                                                                                                                                                                                                                                                                             20-MAY-2002; 2002US-0381857P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 ACTGCACACAGGACCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH65592 standard; DNA; 20
                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                           Nalseth AE;
                                                                                                                                                                                                                                                                                                                   (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-035034/03.
                                                                                                                                                                                                                          WO2003099215-A2.
                                                                                                                                                                                                    Homo sapiens
                                                                                           25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2004
                                                                                                                                                                                                                                                04-DEC-2003
                                                                                                                                                                                                                                                                                                                                            Crosby SD,
                                                                    ADH65235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH65592;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                       RESULT 1680
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Matches
                                   ADH65235
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to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNN sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabeces; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human glucocorticoid receptor-specific antisense oligonucleotide #3155.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 6 A; 8 C; 4 G; 2 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 2426; 985pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1928 ACTGCACACACGACCTGT 1945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19
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                                                                                                                                                    20-MAY-2003; 2003WO-US016084.
                                                                                                                                                                                                                           20-MAY-2002; 2002US-0381857P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 88.9
les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                Crosby SD, Nalseth AE;
                                                                                                                                                                                                                                                                                                     (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHAA ) PHARMACIA CORP
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WO2003099215-A2
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invention also relates to a method of identifying rice plants and seeds deficient in the Rf-1 gene using Rf-1 gene-related sequences; and a method for the regulation of Rf-1-mediated fertility recovery using Rf-1 gene antisense oligonucleotides. The method of the invention restores pollen fertility suppressed by BT cytoplasmic male sterility in rice, which is useful for seed production and for investigation of the mechanisms involved in sterility. Sequences ADI09991-ADI0998 and ADI09999-ADI10038 represent PCR primers used in an example from the invention in RFLP (restriction fragment length polymorphism) or PCR analysis of rice chromosome 10 to determine markers.

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Gaps

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Indels

0; Mismatches

Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;

Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

0.48;

88.98;

Local Similarity 88.5 Les 16; Conservative

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Query Match
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Matches
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                                                                                                                                                        to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, present DNA sequence repersents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of restoring fertility to rice with BT cytoplasmic male sterility. The invention involves transforming rice with a nucleic acid encoding either the rice fertility restores protein Rf-1 ADI10045 or a protein at least 70% homologous to rice Rf-1. Specifically, the Rf-1 protein-encoding nucleic acids used in the method of the invention are ADI09997, ADI10039-ADI10043 or ADI10050-ADI10055. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rf-1 rice genomic sequences on chromosome 10 for fertility restoration to rice BT-type male sterility cytoplasm.
                                           New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                            The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rice; plant; fertility restoration; cytoplasmic male sterility; BT; fertility restorer; Rf-1; chromosome 10; transgenic plant; identification; regulation; pollen fertility; seed production; marker; RFLP; restriction fragment length polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rice chromosome 10 RFLP marker G4003 HindIII PCR primer, SEQ ID NO:3
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                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kuraya Y;
                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 6 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suzuki S,
                                                                                                            Claim 4; SEQ ID NO 3155; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; SEQ ID NO 3; 261pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                        1928 ACTGCACACACGACCTGT 1945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hiei Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                        18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-MAR-2003; 2003WO-JP003154.
                                                                                                                                                                                                                                                                                                                                                   88.98;
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                                                                                                                                                                                                                                                                                                                                                                                                                            ACTGCACACAGGACCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI09973 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NISB ) JAPAN TOBACCO INC. (SYGN ) SYNGENTA LTD.
                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Takakura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-108821/11.
                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI09973;
                                                                                                                                                                                                                                                                                                                                         Query Match
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The invention relates to an antisense oligonucleotide targeted to a nucleic acid encoding the human PPP3CB polypeptide and inhibits expression of the PPP3CB polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, i.e. a phosphorothioate linkage, at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense oligonucleotides are useful for inhibiting expression of the PPP3CB polypeptide and in preparation of a composition for treating autoimmune disorders or Alzheimer's disease. This sequence represents an antisense oligonucleotide target region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding PPP3CB, useful for treating an autoimmune disorder,
                                                                                                                                                                                                                                                                    Human, PPP3CB; as; antisense oligonucleotide; phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; autoimmune disorder; Alzheimer's disease; immunosuppressive; nootropic; neuroprotective.
                                                                                                                                                                                                                                    Human PPP3CB DNA antisense oligonucleotide target region #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4 A; 5 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; SEQ ID NO 86; 91pp; English.
2044 ACCGACGAGTACCTGGAC 2061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dobie KW
                                13
                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-JUL-2002; 2002US-00210723.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-JUL-2002; 2002US-00210723.
                                  2 ATCGACGAGTACCTGAAC
                                                                                                                            ADJ53550 standard; DNA; 20
                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-142663/14.
                                                                                                                                                                                                                                                                                                                                                                                    US2004023382-A1
                                                                                                                                                                                                                                                                                                                                                  sapiens.
                                                                                                                                                                                                  06-MAY-2004
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                                                                                                                                                                ADJ53550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dean NM,
                                                                                     RESULT 1684
ADJ53550/c
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0.4%; Score 14.8; DB 1; Length 20;

Query Match

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ADJ53478

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WPI; 2004-203534/19.
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es 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                    05-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an antisense oligonucleotide targeted to a nucleic acid encoding the human PPP3CB polypeptide and inhibits expression of the PPP3CB polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, i.e. a phosphorothicate linkage, at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense oligonucleotides are useful for inhibiting expression of the PPP3CB polypeptide and in preparation of a composition for treating autoimmune disorders or Alzheimer's disease. This sequence represents an antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding PPP3CB, useful for treating an autoimmune disorder,
                                                                                                                                                                                                                                                                                                                                                                    Human, PPP3CB; ss; antisense oligonucleotide; phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; autoimmune disorder; Alzheimer's disease; immunosuppressive; nootropic; neuroprotective.
                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Pred. No. 1.7e+03;
                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
Pred. No. 1.7e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                Human PPP3CB DNA antisense oligonucleotide #1.
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                                                           2129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
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                                                                                                                                                                                                      ADJ53478 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-JUL-2002; 2002US-00210723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JUL-2002; 2002US-00210723
                                                             2112 CTCCAGCTCCTCAGGGGA
                                                                                                      CTCCAGCTCCTCGGGTGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADJ60733 standard; DNA; 20
                                                                                                                                                                                                                                                                                        (first entry)
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Similarity 88.5
16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-142663/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2004023382-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                        06-MAY-2004
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                                                                                                  18
                                                                                                                                                                                                                                                 ADJ53478;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dean NM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ60733;
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Best Local
Matches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1686
ADJ60733
ID ADJ60733
XX
AC ADJ60733
XX
DT 06-MAY-2
                                                                                                                                                                RESULT 1685
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The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene (s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g.,
                                                     interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes (CR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin, IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1; Length
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 6 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide associated to Tryptase-a #77.
               Oligonucleotide associated to PDE4A #16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 1589; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A,
H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1573 CAGGTGGCCCGGGCATG 1590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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                                                                                                                                                                                                                                                                                                         25-JUL-2003; 2003WO-US023509.
                                                                                                                                                                                                                                                                                                                                                  29-JUL-2002; 2002US-0399076P.
                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC.
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Matches

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Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the cilgonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease. Copen chosen from airway inflammation, allergy(ies), asthma, impeded copen airway inflammation, allergy(ies), asthma, impeded copen copen in the present sequence respiratory distress syndrome (COPD), allergic rhinitis (AR), acute respiratory distress syndrome coherunction. The present sequence represents an oligonucleotide of the invantion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                      genes e.g.,
cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; G protein-coupled receptor 6; GPCR-6; ss;
antisense oligonucleotide; phosphorothicate linkage;
2'-O-methoxyethyl sugar moiety; 5-methylcytosine; metabolic disorder;
aberrant signal transduction; brain tissue; neuronal disorder;
motor disorder; sensory disorder; psychiatric disorder;
behavioural disorder; drug addiction; chemical addiction; neuroleptic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human G protein-coupled receptor 6 DNA antisense oligonucleotide #39
                                                                                                                                                                                                                                                                                                   initiation codons and introns of respiratory disease-relevant gener
CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
disease e.g., asthma.
                                                                                                                                                                                                                                                                                        Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indel8
                                                                                                                                                                                                                     Miller S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 10 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                     Aguilar D,
                                                                                                                                                                                                                                                                                                                                                             Claim 2; SEQ ID NO 1397; 85pp; English.
                                                                                                                                                                                                                      Tang L, Sandrasagra A,
in S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3196 CCGGAGCTGGAGGATCCC 3213
                                                                                                                                                                                                                                   Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
                                                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                      25-JUL-2003; 2003WO-US023509
                                                                                                                                                                 29-JUL-2002; 2002US-0399076P
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                             WPI; 2004-203534/19.
                                                                                WO2004011613-A2
                                                                                                                                                                                                                                  Shahabuddin S,
                                                       Homo sapiens
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                                                                                                           05-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
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The invention relates to an antisense oligonucleotide targeted to a nucleic acid encoding the human G protein-coupled receptor 6 (GPCR-6), which specifically hybridises with the nucleic acid encoding the GPCR-6 and inhibits expression of the GPCR-6. The antisense oligonucleotide comprises at least one modified internucleoside linkage, i.e. a phosphorothicate linkage, at least one modified sugar moiety, preferably a 2-0-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense oligonucleotides are useful for inhibiting expression of the GPCR-6 and in preparation of a composition for treating a disease or condition associated with GPCR-6. e.g., a metabolic disorder, aberrant should associated with GPCR-6, a neuronal, motor, sensory, systhiatric or behavioural disorder or drug or chemical addiction. This sequence represents an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense compound targeted to nucleic acids encoding G protein-coupled receptor 6 (GPCR-6), useful for treating animal having disease associated with GPCR-6 e.g. metabolic, neuronal, motor, sensory or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; G protein-coupled receptor 6; GPCR-6; ss; antisense oligonucleotide; phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; metabolic disorder; aberrant signal transduction; Drain tissue; neuronal disorder; motor disorder; sensory disorder; psychiatric disorder; behavioural disorder; drug addiction; chemical addiction; neuroleptic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
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                                                                                                                                        31-JUL-2002; 2002US-00210479.
                                                                                                                                                                                                           31-JUL-2002; 2002US-00210479
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.9'
                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          behavioral disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-142661/14.
US2004023380-A1
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                                                                      05-FEB-2004
                                                                                                                                                                                                                                                                                                                                                       Monia BP,
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16; Conservative
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                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                    Dobie KW;
                                                                                                                                          WPI; 2004-142660/14.
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Best Local Similarity
US2004023379-A1
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modified_base
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                                                                                                                    Bennett CF,
                         05-FEB-2004
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                                                                                                                                                                                                   disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                              nucleic acid encoding the human G protein-coupled receptor 6 (GPCR-6), which specifically hybridises with the nucleic acid encoding the GPCR-6 and inhibits expression of the GPCR-6. The antisense oligonucleotide comprises at least one modified internucleoside linkage, i.e. a phosphorothicate linkage, at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense oligonucleotides are useful for inhibiting expression of the GPCR-6 and in preparation of a composition for treating a disease or condition associated with GPCR-6, e.g., a metabolic disorder, aberrant signal transduction in brain tissue, a neuronal, motor, sensory, psychiatric or behavioural disorder or drug or chemical addiction. This sequence represents an antisense or drug oligonucleotide target region of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; endocrine; hepatoma-derived growth factor inhibitor; hepatoma-derived growth factor; metabolic disorder; hyperproliferative; human; 88; antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                      Novel antisense compound targeted to nucleic acids encoding G protein-coupled receptor 6 (GPCR-6), useful for treating animal having disease associated with GPCR-6 e.g. metabolic, neuronal, motor, sensory or behavioral disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "OTHER= 2'-0-Methoxyethyl (2'-MOE) nucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                     invention relates to an antisense oligonucleotide targeted to a
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatoma-derived growth factor antisense oligo seqid 127.
                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                              Example 15; SEQ ID NO 112; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                    2015 ACCTGGACCGTGTCCTTA 2032
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/mod_base= OTHER
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            31-JUL-2002; 2002US-00210479.
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nes 16; Conservative
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/*tag=
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                                 (ISIS-) ISIS PHARM INC
                                                          Dobie KW;
                                                                                 WPI; 2004-142661/14.
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                                                          Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1690
ADJ45357/c
                                                                                                                                                                                                                                                                                                                                                                                                  Best Loc
Matches
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The invention describes a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding hepatoma-derived growth factor, and inhibits the expression of hepatoma-derived growth factor. The compound, composition and methods are useful for treating a disease or condition associated with hepatoma-derived growth factor, such as a metabolic disorder, or a hyperproliferative disorder, e.g. cancer, which is selected from hepatoma, leiomyoma, esophageal cancer or ovarian cancer. They are also useful in research and diagnostics for modulating the expression of hepatoma-derived growth factor. This sequence represents a human hepatoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytostatic; endocrine; hepatoma-derived growth factor inhibitor; hepatoma-derived growth factor; metabolic disorder; hyperproliferative; human; ss; antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                              2
                                                                                                                                                                                                                                                                                                                                                                       New compound, particularly an antisense oligonucleotide targeted to
nucleic acid encoding a hepatoma-derived growth factor, useful for
treating a hyperproliferative disorder e.g. cancer, or a metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
0
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatoma-derived growth factor antisense oligo segid 56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 8 A; 1 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             derived growth factor antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15; SEQ ID NO 127; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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/*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= c
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ45286 standard; DNA; 20 BP.
31-JUL-2002; 2002US-00210429
                                                                         31-JUL-2002; 2002US-00210429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 ccrrrrcrrcrrrarc
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The present invention relates to antisense compounds, compositions and methods for modulating the expression of Centromere protein B. The compound, composition and methods are useful for treating diseases or conditions associated with Centromere protein B, such as hyperproliferative disorders (e.g. cancer), autoimmune disorders e.g. rhematoind arthritis, scleroderma, Raymaud's syndrome or systemic lupus erythematosus. The invention is also useful in gene therapy. The present sequence is human Centromere protein B antisense oligonucleotide used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorothioate; 2' MOE; breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; all stone; triglyceridaemia; obesity; hepatitis; hepaticellular carcinoma; aromatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antiinflammatory; virucidal:
                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotide targeted to a nucleic acid encoding
Centromere protein B, useful for treating a disease, e.g. cancer,
rheumatoid arthritis, scleroderma, Raynaud's syndrome or systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense DNA oligo used to modulate human LRH1 expression SeqID 3016.
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                             16. .20
/*tag= c
/mod_bag= c
/mod_e= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88; liver related homologue-1; LRH1; NR5A2; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 5 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; SEQ ID NO 17; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1262 AGGACCGGCCGCCAAGC 1279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AGGACTGGGCAGCCAAGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
                                                                                                                                                                                  18-JUN-2002; 2002US-00176277.
                                                                                                                                                                                                                   18-JUN-2002; 2002US-00176277.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ18466 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAY-2004 (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                     WPI; 2004-052175/05
                                                                                                                   US2003232443-A1
                                  modified base
                                                                                                                                                                                                                                                                                                                                                                                                       erythematosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified base
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                                                                                                                                                                                                                                                                                    Bennett CF,
                                                                                                                                                   18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ18466;
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                                                                                                                                                                                                                                                                                                                                                                                                  to, and which specifically hybridises with a nucleic acid molecule encoding hepatoma-derived growth factor, and inhibits the expression of hepatoma-derived growth factor. The compound, composition and methods are useful for treating a disease or condition associated with hepatoma-derived growth factor, such as a metabolic disorder, or a hyperproliferative disorder, e.g. cancer, which is selected from hepatoma, leiomyoma, esophageal cancer or ovarian cancer. They are also useful in research and diagnostics for modulating the expression of hepatoma-derived growth factor. This sequence represents a human hepatoma-derived growth factor. This sequence represents a human hepatoma-derived growth factor antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Phosphorothioate backbone where all cytidines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Centromere protein B; hyperproliferative disorder; cancer; autoimmune disorder; rheumatoid arthritis; scleroderma; Raynaud's syndrome; systemic lupus erythematosus; gene therapy; cytostatic; immunosuppressive; dermatological; antiinflammatory; human; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                    The invention describes a compound 8-80 nucleobases in length targeted
 /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                   New compound, particularly an antisense oligonucleotide targeted to a nucleic acid encoding a hepatoma-derived growth factor, useful for treating a hyperproliferative disorder e.g. cancer, or a metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Centromere protein B antisense oligonucleotide, ISIS #156869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 2 A; 9 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 56; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5-methylcytidines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        920 GCTTCTTCCTGTTCATCC 937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
                                                                                                   31-JUL-2002; 2002US-00210429
                                                                                                                                    31-JUL-2002; 2002US-00210429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCTCTTCCTCTTCATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ26848 standard; DNA;
                                                                                                                                                                    (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                     Bennett CF, Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                    WPI; 2004-142660/14.
                                    US2004023379-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-MAY-2004
                                                                  05-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                      disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ26848;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
Matches
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Gaps .; 0 vivlemore401-10.rng

phosphorothioate; 2' MOE; breast cancer; dyalipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis; hepatocallular carcinoma; aromatase; cytostatic; antilipaemic; antiarteriosclerotic; anorectic; hepatotropic; litholytic; antinflammatory; virucidal.

Antisense DNA oligo used to modulate human LRH1 expression SeqID 2243 ss; liver related homologue-1; LRH1; NR5A2; antisense; nucleotides. All

/note= "OTHER= 2' methoxyethyl (2' MOE) nuc cytidine nucleobases are 5-methylcytidine."

/*tag= a /mod_base= OTHER /note= "OTHER= 2'

modified base

'label = OTHER = phosphorothioate backbone

base= OTHER

Ω

/*tag=

Key modified_base

sapiens

Homo

Synthetic.

socation/Qualifiers

nucleotides. All

/mod_base= OTHER /note= "OTHER= 2' methoxyethyl (2' MOE) nu cytidine nucleobases are 5-methylcytidine.

*tag=

modified base

01-JUL-2003; 2003WO-US020865. 01-JUL-2002; 2002US-0392813P.

WO2004003201-A2.

08-JAN-2004.

(PHAA) PHARMACIA CORP

WPI; 2004-083058/08.

Kane CD;

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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants therefor Specifically, it refers to compositions 8-30 nucleobases in clength that target a portion of an active site on the nucleic acid molecule encoding LRH1 (also known as NR5A2). LRH1 is a monomeric orphan unclear receptor protein that functions as a tissue specific or transcription factor. The present invention describes antisense transcription factor. The present invention describes antisense oligonucleotides that comprise at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MDE) and at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MDE) and at least one modified sugar moiety, a 2'-0-methoxyethyl (2' MDE) and at least one modified succer, diagnosing a disease associated with LRH1, such as best cancer, dyslipidaemia, atherosclerosis, low HDL (high density lipoprotein), hypercholesterolaemia, gall stones, triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit characteric, antilipaemic, antiarteriosclerotic, anorectic, hepatotropic, litholytic, antilipaemic, and virucidal activities. This cliconucleotide sequence is an antienness and or a condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                          /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                                                                                                                                                                                                                  /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All cytidine nucleobases are 5-methylcytidine."
                        /label= OTHER= phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 13 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 15; SEQ ID NO 3016; 909pp; English.
                                                                                    /mod_base= OTHER
/note= "OTHER= 2'
                                                                                                                                                                             /*tag= c
/mod_base= OTHER
'mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2003; 2003WO-US020865.
                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-2002; 2002US-0392813P.
                                                                    ๙
                                                                *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-083058/08.
                                                                                                                                                                                                                                                                                 WO2004003201-A2
                                            modified base
                                                                                                                                                    modified base
                                                                                                                                                                                                                                                                                                                           08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kane CD;
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This invention relates to novel antisense compounds useful for modulating the expression of liver related homologue-1 (LRH1) and splice variants thereof. Specifically, it refers to compositions 8-30 nucleobases in length that target a portion of an active site on the nucleic acid molecule encoding LRH1 (also known as NB5A2). LRH1 is a monomeric orphan nuclear receptor protein that functions as a tissue specific cranscription factor. The present invention describes antisense coligomucleotides that comprise at least one modified internucleoside linkage, a phosphorothicate linkage; at least one modified sugar moiety, a 2-0-methoxyethy1 (2' MDE) and at least one modified nucleobase, a 5-comethyquidine. These antisense compounds are useful for treating or diagnosing a disease associated with LRH1, such as breast cancer, dyslipidaemia, atherosclerosis, low HDL (high density lipoprotein), high continued acide acide or chronic triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with their anomates and continue exhibit the continued and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic, antilipaemic, antiarteriosclerotic, anorectic, hepatotropic, litholytic, antiinflammatory and virucidal activities. This oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRHI protein of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
0
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     318 CCCCACTCCCTCCATCTC 335
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Matches 16; Conservative
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ADJ17693 standard; DNA; 20

Query Match

g 8

(first entry)

20-MAY-2004

ADJ17693;

RESULT 1694
ADJ17693
ID ADJ1769:
XX
AC ADJ1769:
XX
DT 20-MAY-:

Sequence 20 BP; 3 A; 12 C; 0 G; 5 T; 0 U; 0 Other;

New antisense oligonucleotides targeted to a nucleic acid encoding liver related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.

Example 15; SEQ ID NO 2243; 909pp; English

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20-MAY-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to the primary culture of adipocytes for gene therapy that stably carry a foreign secreted protein. Further disclosed is a method for producing the adipocytes, a transplant composition for gene therapy containing the cells, and a method for gene therapy and animals with the transplanted adipocytes. The proteins are preferably insulin or glucagon-like peptide 1. The cells of the invention are useful for gene therapy, particularly as a substitute for bone marrow cells or hepatocytes. The primarily cultured adipocytes are useful for ex vivo gene therapy. The current sequence represents a primer used in an example
                                                                                                                                                                                                         Gene therapy, adipocyte, secreted protein; insulin;
glucagon-like peptide 1; bone marrow cell; hepatocyte; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                Primary culture of adipocytes for gene therapy useful in treatment of diabetes.
                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
        Length 20;
                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;
       Score 14.8; DB 1;
Pred. No. 1.7e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; SEQ ID NO 8; 68pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2238 CCACCCTGCTGCTGGTGC 2255
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                                                                     2 ccccacrccccaarere 19
                                                                                                                        ADM28947 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                     18-JUN-2003; 2003WO-JP007721.
                                                                                                                                                                                                                                                                                                                          18-JUN-2002; 2002JP-00177648.
19-AUG-2002; 2002JP-00237974.
         0.4%;
                                                  318 CCCCACTCCCTCCATCTC
                                                                                                                                                                (first entry)
       Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                     Primer GLP-5', seq id 8.
                                                                                                                                                                                                                                                                                                                                                        (EISA ) EISAI CO LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                             Ito M, Saito Y;
                                                                                                                                                                                                                                                             WO2003106663-A1.
                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                20-MAY-2004
                                                                                                                                                                                                                                                                                 24-DEC-2003
                                                                                                                                             ADM28947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                     RESULT 1695
ADM28947
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Matches
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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2 MOS wings and a deoxy age. Used during the antisense inhibition of thuman Navl.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; 88.
                                                      Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
Chimeric phosphorothioate oligonucleotide to target Navl.3 #4481.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric phosphorothioate oligonucleotide to target Navl.3 #5791.
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Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 ·U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 4481; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1148 AGCTGCCTGCCGACCCCA 1165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AGCTGCATGCCGACCACA 18
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                                                                                                                                                                                                                                                                                                                                                    14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                                                                                                              14-AUG-2002; 2002US-0403416P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                 WO2004016754-A2
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                                                                                                                                                                             Synthetic.
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ADK77147 standard; DNA; 20

ADK77147 RESULT 1696
ADK77147
ID ADK77147
XX
AC ADK77147
XX

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2001US-0274194P.
2001US-0274281P.
2001US-027432P.
2001US-0275235P.
2001US-0275579P.
2001US-0275579P.
2001US-0275601P.
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2001US-0277239P
2001US-027733P
2001US-027733P
2001US-027731P
2001US-027731P
2001US-0278132P
2001US-0278994P
2001US-0278999P
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2001US-0303230P.
2001US-0303237P.
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                                                                   07-MAR-2002; 2002US-00094886
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2001US-0280802P.
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2001US-0312916P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-0311978P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-0313182P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-AUG-2001; 2001US-0313626P
21-AUG-2001; 2001US-0314018P
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PATTURAJAN M.
BURGESS C E.
VERNET C A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GORMAN L.
MALYANKAR U M.
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PADIGARU M.
TAUPIER R J.
MILLER C E.
CASMA C E A.
GANGOLLI E A
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SMITHSON G.
   US2004002120-A1
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                                    01-JAN-2004
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(VERN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PADI/)
(TAUP/)
(MILL/)
(CASM/)
(PENA/)
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(GORM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SHEN/
   The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate beadache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy, or ataxia. The present but not limited a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; NOVX; PCR; ss; G protein-coupled receptor; GPCR; cardiomyopathy; atherosclerosis; hypertension; congenital heart defect; aortic stenosis; atrial septal defect; ASD; atrioventricular canal defect; ductus arteriosus; pulmonary stenosis; subaortic stenosis; ventricular septal defect; VSD; tuberous sclerosis; scleroderma; obesity; adrenoleukodystrophy; congenital adrenal hyperplasia; prostate cancer; neoplasm; adenocarcinoma; lymphoma; uterine cancer; haemophilia;
                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding aball.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Albright's heraditary osteodystrophy; diabetes; infectious diseases; anorexia; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; haematopoietic disorder; metabolic disorder; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypercoagulability; idiopathic thrombocytopenia purpura; immunodeficiency; graft-versus-host disease; AIDS; bronchial asthma; Crohn's disease; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 0.4%; Score 14.8; DB 1; Length 20; Local Similarity 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 6 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 5791; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1147 GAGCTGCCTGCCGACCCC 1164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAGCTGCATGCCGACCAC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADL24187 standard; DNA; 20 BP
                                                               14-AUG-2003; 2003WO-US025465.
                                                                                                  14-AUG-2002; 2002US-0403416P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16; Conservative
                                                                                                                               (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                WPI; 2004-203785/19.
WO2004016754-A2
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                                                                                                                                                                 Roberds SL;
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셤 ð

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Kekuda R, Tchernev VT, Liu X, Spytek KA, Patturajan M;
Burgess CE, Vernet CAM, Li L, Gorman L, Malyankar UM, Boldog FL;
Guo X, Shenoy SG, Padigaru M, Taupier RJ, Miller CE, Casman SJ;
Pena CEA, Gangolli EA, Gusev VY, Smithson G, Zerhusen BD, Gerlach V;
Pochart PP, Fernandes ER, Shinkets RA, Rastelli L, Spaderna SK;
Larochelle WJ, Zhong M, Khramtsov NV, Voss EZ, Herrmann JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated G protein-coupled receptor polypeptides, referred as NOVX, useful for treating scleroderma, obesity, congenital adrenal hyperplasia, prostate cancer, hemophilia, AIDS, bronchial asthma, Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          related) polypeptides (designated NOVX) and the polypubleotides encoding them. The polypeptides and polypubleotides are useful as therapeutics in them manufacture of medicaments for treating syndromes associated with human diseases. The sequences are useful for treating a disorder associated with aberrant NOVX expression or activity such as cardiomyopathy, atherosclerosis, hypertension, congenital heart defects, eartis septal septal defect (ASD), atrioventricular canal defect, ductus arteriosus, pulmonary stenosis, subsoctic stenosis, ventricular septal defect (VSD), tuberous sclerosis, scleroderma, obesity, adrenoleukodystrophy, congenital adrenal hyperplasia, prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer, neoplasm, adenocarcinoma, lymphoma, uterine cancer, haemophilia, hypercoagulability, idiopathic thrombocytopenia purpura, immunodeficiencies, graft-versus-host disease, AIDS, bronchial asthma, Crohn's disease, multiple sclerosis, Albright's hereditary osteodystrophy, diabetes, infectious diseases, anorexia, nourodegenerative disorders, Albrightes's disease, Parkinson's disease, neurodegenerative disorders, Albrightes's disease, Parkinson's disease, immune disorders, haematopoietic disorders and metabolic disorders. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence represents a PCR primer used in analysis of human NOVX sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to human G protein-coupled receptor-related (GPCR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA array; support; functional group; mismatch detection; virus identification; bacterium identification; p53; PCR; primer; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.76+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 47; SEQ ID NO 232; 287pp; English.
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es 16; Conservative
                                                                                                                                                        ZHONG M.
KHRAMTSOV N V.
VOSS E Z.
                                                                                                                 SPADERNA S K.
LAROCHELLE W J.
                    GERLACH V.
POCHART P F.
FERNANDES E R.
                                                                                SHIMKETS R A.
                                                                                                                                                                                                                    HERRMANN J L.
ZERHUSEN B D.
                                                                                                   RASTELLI L.
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-212692/20
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Best Local S
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(KHRA/)
(VOSS/)
(HERR/)
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                                                                                                   RAST/)
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                                                                              SHIM/
                                                           FERN/
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                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method of producing a cDNA array on a support, which involves introducing a functional group to one edge part of 2 or more types of single stranded cDNA (known sequence) for fixing to a support and combining each strand of cDNAs with a support through a functional group for binding such that each strand of cDNA is mutually isolated and fixed. The method is useful for preparing a cDNA array, which is useful for detecting mismatches, or for identifying (for example) viruses or bacteriar The present sequence is that of a PCR primer which was used for amplification of a region of the human p53 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOVX; antiarteriosclerotic; cytostatic; antidiabetic; antiparkinsonian; neuroprotective; nootropic; antiasthmatic; antiallergic; dimmunosuppressive; antiarthritic; antirheumatic; osteopathic; dermatological; antiinflammatory; anti-HIV; hypotensive; haemostatic; anorectic; gastrointestinal-den; antiuncer; antimicrobial; antipsoriatic; neuroleptic; antidepressant; anabolic; eating disorders-Gen; antidietrility; nephrotropic; gene therapy; antisense gene therapy; human disease; atherosclerosis; cancer; diabetes; Alzheimer's disease; Parkinson's disease; allergy; immune disease; graft-versus-host disease; osteoarthritis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             depression; anorexia; fertility; glomerulonephritis; chromosome mapping; tissue typing; PCR; primer; ss; human.
                                                                                                                                                                                                                                                                            Producing cDNA array on support with introduction of the coupling group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       scleroderma; systemic lupus erythematosus; AIDS; hypertension; haemophilia; idiopathic thrombocytopenic purpura; obesity; inflammatory bowel disease; Crohn's disease; ulcerative colitis; infectious disease; psoriasis; multiple sclerosis; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human NOVX protein-related PCR primer SeqID133.
                                                                                                                                                                                                                                                                                                 using a PCR primer containing the group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID NO 6; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM74292 standard; DNA; 20 BP.
                                                                                                                 06-AUG-2002; 2002JP-00228971
                                                                                                                                                           06-AUG-2002; 2002JP-00228971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2695 CCACTTCCCACCCTGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 ccacrreccacerecae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                     WPI; 2004-308703/29
                                                                                                                                                                                                (CANO ) CANON KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2004015079-A2
                                       JP2004069488-A.
  Homo sapiens.
                                                                              04-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM74292;
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This invention relates to novel human NOVX proteins and the DNA sequences which encode them. The invention may be useful for the development of compounds with an antiarteriosclerotic, cytostatic, antidiabetic, compounds with an antiarteriosclerotic, cytostatic, antidiabetic, compounds with an antiarteriosclerotic, cytostatic, antidiabetic, costeopathic, dermatological, antidifammatory, antidiathematic, costeopathic, dermatological, antidifammatory, antidepressave, antianthritic, antidicerility or nephrotropic activity. In addition, the sequences den, antidifertility or nephrotropic activity. In addition, the sequences of inclosed may be useful for gene therapy or antisense gene therapy. The NOVX polypeptides, nucleic acid and antibody are useful for manufacturing a medicament for treating a syndrome associated with a human disease. The NOVX polypeptides and polymucleotides are also useful for diagnosing, treating and preventing disorders associated with aberrant expression or aberrant physicological interactions of the polypeptide, for example of insease, asthma, allergies, immune disease, graft-versus-host disease, osteoparthritis, theumatoid arthritis, selected arthritis, selected arthritis, selected arthritis, infectious disease, parkinson's crivematosus, AIDS, hypertension, heamophilia, idiopathic cup a disease cupromatory burner, obesity, inflammatory bowel disease, contextissis, multiple selected with altered levels of the NOVX polypeptides and nucleic acid molecules or the nucleic control acid molecule or for screening for molecules that inhibit or enhance control selected with altered levels of the NoVX polypeptide or the nucleic control second molecules that inhibit or enhance control second molecules that inhibit or an edicine, or pharmacogenomics. The present sequence is that of a pcR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel polypeptides (NOVX) and nucleic acid molecules, useful for diagnosing, preventing or treating NOVX-associated disorders, e.g. atherosclerosis, cancer, diabetes, Alzheimer's disease, Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example C; SEQ ID NO 133; 233pp; English.
                                                                            09-AUG-2002; 2002US-0402205P.

13-AUG-2002; 2002US-0402205P.

13-AUG-2002; 2002US-0403517P.

13-AUG-2002; 2002US-040354P.

15-AUG-2002; 2002US-040369FP.

26-AUG-2002; 2002US-0406318P.

27-AUG-2002; 2002US-0406302P.

06-SEP-2002; 2002US-00236392.

13-SEP-2002; 2002US-00236392.

01-NOV-2002; 2002US-0024943.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease, asthma or allergies.
                      07-AUG-2003; 2003WO-US024931
                                                                                                                                                                                                                                                                                                     06-AUG-2003; 2003US-00635149
                                                                                                                                                                                                                                                                                                                                                                                   . W, Guo X,
Anderson DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                Edinger SR;
                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-180660/17.
                                                                                                                                                                                                                                                                                                                                                                                       Ji W,
                                                                                                                                                                                                                                                                                                                                                                                                              ₹,
                                                                                                                                                                                                                                                                                                                                                                                                                                Stone DJ,
                                                                                                                                                                                                                                                                                                                                                                                       Zhong M,
                                                                                                                                                                                                                                                                                                                                                                                                           Spytek
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Match 0.4%; Score 14.8; DB 1; Length 20; Local Similarity 88.9%; Pred. No. 1.7e+03; es 16; Conservative 0; Mismatches 2; Indels Sequence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other; Query Match

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Gaps

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1512
                           20
1495 GGCCTGGACTACTCCTTC
                         3 GGCCTGGACTGCTTCTTC
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g

Matches

hyperproliferative disease; cell cycle progression; protein phosphorylation; tumour growth; cancer; apoptosis; protein cancer; inflammation; fibrosis; fibrotic disease; scarring; peritoneal adhesion; lung fibrosis; conjunctival scarring; cytostatic; antiinflammatory; vulnerary; ss. Human; Jun N-terminal kinase; JNK; Jun N-terminal kinase 2; JNK2; Human Jun N-terminal kinase 2 (JNK2) oligonucleotide #3. ADN48320 standard; DNA; 20 BP. 07-AUG-1998; 98US-00130616. 07-ARR-1999; 99US-00287796. 11-SEP-1999; 99US-00396902. 31-JAN-2001; 2001US-00774809. 15-JAN-2003; 2003US-00345444 (first entry) DEAN N M.
MONIA B P.
NERO P S.
GAARDE W A. US2004029823-A1. MCKAY R. Homo sapiens. 01-JUL-2004 13-AUG-1997; 12-FEB-2004 ADN48320; (MCKA/) (DEAN/)

Malcolm R; Macdougall JR;

Padigaru M, Catterton E,

Rieger DK, Gorman L,

Gaarde WA; Nero PS, Dean NM, Monia BP, Mckay R,

(MONI/) N (NERO/) N (GAAR/) C

New oligonucleotides, which specifically hybridizes with Jun N-terminal kinase protein, useful in diagnosing or treating inflammation, fibrosis or a fibrotic or hyperproliferative disease or condition. WPI; 2004-168941/16.

Claim 25; SEQ ID NO 31; 71pp; English.

The invention relates to an oligonucleotide comprising 8-30 nucleotides connected by covalent linkages, where the oligonucleotide has a sequence specifically typridisable with a nucleic acid encoding a Jun N-terminal kinase (JNK) protein and modulates the expression of the JNK protein. The invention also relates to a pharmaceutical composition comprising the cligonucleotide(s) or its bioequivalent and a pharmaceutical carrier, a method of treating an animal having, suspected of having or prone to having a hyperproliferative disease, a method of modulating the expression of a JNK protein in cells or tissues, a method of modulating cell cycle progression, phosphorylation of a protein that promotes one or JNK protein and expression of a cellular protein that promotes one or one metastatic events in cultured cells or the cells of an animal, a method of inducing apoptosis in a cell, a method of treating a mainal a cell, a method of treating a having a completion of treating an animal having a cumple of treating an animal having a condition associated with a JNK protein. The oligonucleotide and composition are useful in diagnosismy or treating a disease or condition characterised by a creduction in apoptosis (e.g. prostate cancer), a disease or condition associated disease or condition (e.g. scarring, peritoneal adhesions, lung fibrosis, conjunctival scarring) or a hyperproliferative disease or condition (e.g. cancer), or in inhibiting the growth of a tumour. This sequence

Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 U; 0 Other;

vivlemore401-10.rng

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ADN35260/
8888888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an oligonucleotide comprising 8-30 nucleotides connected by covalent linkages, where the oligonucleotide has a sequence specifically hybridiashle with a nucleic acid encoding a Jun N-terminal kinase (JNK) protein and modulates the expression of the JNK protein. The invention also relates to a pharmaceutical composition comprising the oligonucleotide(8) or its bioequivalent and a pharmaceutical carrier, a method of treating an animal having, suspected of having or prone to having a hyperproliferative disease, a method of modulating the expression of a JNK protein in cells or tissues, a method of modulating cell cycle progression, phosphorylation of a protein phosphorylated by a compose more metastatic events in cultured cells or the cells of an animal, a method of inhibiting the growth of a tumour in an animal, a method of indivining the growth of a tumour in an animal, a method of indivinion characterised by a ceduction in apoptosis and a disease or condition characterised by a disease or condition associated
                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotides, which specifically hybridizes with Jun N-terminal kinase protein, useful in diagnosing or treating inflammation, fibrosis or a fibrotic or hyperproliferative disease or condition.
                                                                                                                                                                                                                                                             hyperproliferative disease; cell cycle progression; protein phosphorylation; tumour growth; cancer; apoptosis; prostate cancer; inflammation; fibrosis; fibrotic disease; scarring; peritoneal adhesion; lung fibrosis; conjunctival scarring; cytostatic;
                                    Gaps
                                                                                                                                                                                                                                                Human; Jun N-terminal kinase; JNK; Jun N-terminal kinase 2; JNK2;
                                    ö
            Length 20;
                                                                                                                                                                                                                         Human Jun N-terminal kinase 2 (JNK2) oligonucleotide #14.
                                    2; Indels
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           Score 14.8; DB 1;
Pred. No. 1.7e+03;
0; Mismatches 2;
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                                                            1695
                                                                                                                                                                                                                                                                                                               antiinflammatory; vulnerary; ss.
                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-1998; 98US-00130616.
07-APR-1999; 99US-00287796.
15-SEP-1999; 99US-00774809.
31-JAN-2001; 2001US-00774809.
                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-2003; 2003US-00345444
                                                                                                                                                                                                                                                                                                                                                                                                                                        97US-00910629
           ch 0.4%;
1 Similarity 88.9%;
16; Conservative
                                                           1678 GACTTCGGGCTGGCCCGG
                                                                                  20 GACTTTGGCCTGGCCCGG
                                                                                                                                                ADN48331 standard; DNA; 20
                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEAN N M.
MONIA B P.
NERO P S.
GAARDE W A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-168941/16
           Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MCKAY R.
                                                                                                                                                                                                                                                                                                                                                               US2004029823-A1.
                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                       12-FEB-2004.
                                                                                                                                                                         ADN48331;
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(GAAR/) (
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(MONI/)
                                                                                                                         RESULT 1702
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with a JNK protein. The oligonucleotide and composition are useful in diagnosing or treating a disease or condition characterised by a reduction in apoptosis (e.g. prostate cancer), a disease or condition associated with a JNK protein (e.g. inflammation, fibrosis), a fibrosis, disease or condition (e.g. scarring, peritoneal adhesions, lung fibrosis, conjunctival scarring) or a hyperproliferative disease or condition (e.g. cancer), or in inhibiting the growth of a tumour. This sequence represents a human JNK2 oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             secondary-ion mass spectrometry; gene analysis; disease diagnosis; species identification; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
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Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 7 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Target sequence of the invention #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hashimoto H;
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28-JUN-2002; 2002JP-00191414.
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                                                                                                                                                                                                                                                                                                                                                                                        0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1678 GACTTCGGGCTGGCCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADN35260 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-203385/19
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Gaps
                                                                                                                                                                                 Human; NOVX; PCR; ss; metabolic disorder; diabetes; obesity; infectious disease; anorexia; cancer; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; immune disorder; haematopoietic disorder; antidiabetic; anorectic; antimicrobial; anabolic; eating disorder; cytostatic; neuroprotective; nootropic; antiparkinsonian; antianaemic; primer.
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2; Indels
0; Mismatches
                    CCACTTCCCACCTGCCC 2712
                                                                                                                                                                                                                                                                                                                                                07-SEP-2001; 2001US-0318120P.
07-SEP-2001; 2001US-0318130P.
07-SEP-2001; 2001US-0318430P.
10-SEP-2001; 2001US-0318430P.
11-SEP-2001; 2001US-0318430P.
11-SEP-2001; 2001US-032816P.
11-SEP-2001; 2001US-032816P.
11-SEP-2001; 2001US-032816P.
12-SEP-2001; 2001US-032816P.
13-SEP-2001; 2001US-032816P.
13-SEP-2001; 2001US-032816P.
13-SEP-2001; 2001US-032816P.
15-SEP-2001; 2001US-032890P.
15-SEP-2001; 2001US-032890P.
15-SEP-2001; 2001US-032890P.
15-SEP-2001; 2001US-032890P.
15-SEP-2002; 2001US-038890P.
15-MAR-2002; 2002US-0366131P.
17-MAY-2002; 2002US-0381664P.
17-MAY-2002; 2002US-0381664P.
17-MAY-2002; 2002US-0381664P.
17-MAY-2002; 2002US-0381664P.
                                       ccactroccaccticac 2
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                                                                                               ADN96339 standard; DNA; 20
                                                                                                                                         (first entry)
                                                                                                                                                             Human NOVX PCR primer #99
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CATTERTON E.
MILLER C E.
SHENOY B G.
PATTURAJAN M.
PENA C E A.
TCHERREN V T.
PADIGARU M.
GUSEV V Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KEKUDA R.
TAUPIER R J.
ANDERSON D W.
VERNET C A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LI L.
GORMAN L.
SPYTEK K A.
                                                                                                                                                                                                                                                                                 US2004067490-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZHONG M.
                                                                                                                                         01-JUL-2004
                                                                                                                                                                                                                                                                                                        08-APR-2004
                    2695
                                                                                                                     ADN96339;
                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (LILL/)
(GORM/)
(SPYT/)
(KEKU/)
(TAUP/)
(VERN/)
(CATT/)
(MILL/)
(SHEN/)
(PATT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TCHE/)
(PADI/)
(GUSE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZHON/)
                                                                         RESULT 1704
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                                                                                     ADN96339
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isolated nucleic acids can be used to express the novel proteins, to detect novel mRNA or a genetic lesion in a novel gene and to modulate its activity. It can also be used in gene therapy for treating or preventing a pathology associated with the protein or nucleic acid. The disorders include metabolic disorders, diabetes, obesity, infectious diseases, anorexia, cancer, neurodegenerative disorders, Alzheimer's disease, Parkinson's disease, immune disorders and haematopoietic disorders. This sequence represents a PCR primer used in analysis of expression of a human NOVX polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to human NOVX polypeptides and polynucleotides. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polypeptide, useful for treating or preventing a pathology associated with the polypeptide, e.g. diabetes, infectious disease, cancer, neurodegenerative disorders or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
                                                                                                                                                                                                                                                                                                                                        Zhong M, Li L, Gorman L, Spytek KA, Kekuda R, Taupier RJ; Anderson DW, Vernet CAM, Catterton B. Miller CE, Shenoy SG; Patturajan M, Pena CEA, Tchernev VT, Padigaru M, Gusev VY; Malyankar UM, Burgess CE, Gerlach V, Casman SJ, Rieger DK; Grosse WM, Smithson G, Peyman JA, Starling G, Rothenberg ME, Larochelle WJ, Shimkets RA, Crabtree J, Raatelli L, Voss EZ; Boldog FL, Edinger SR, Millet I, Macdougall JR, Ellerman K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1343.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1495 GGCCTGGACTACTCCTTC 1512
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Best Local Similarity 88.9%;
Matches 16; Conservative (
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                                                                                                                                      ROTHENBERG M E.
LAROCHELLE W J.
SHIMKETS R A.
CRABTREE J.
RASTELLI L.
VOSS E Z.
BOLDOG F L.
EDINGER S R.
MILLET I.
                                                                                                                                                                                                                                                                            MACDOUGALL J R
           BURGESS C E.
GERLACH V.
CASMAN S J.
RIEGER D K.
GROSSE W M.
SMITHSON G.
PEYMAN J A.
STARLING G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-355290/33.
                                                                                                                                                                                                                                                                                                              CHAPOVAL A.
                                                                                                                                                                                                                                                                                              ELLERMAN K.
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(VOSS/)
(BOLD/)
(EDIN/)
(MILL/)
(MACD/)
(ELLE/)
            (BURG/)
(GERL/)
(CASM/)
(RIEG/)
(GROS/)
(SMIT/)
                                                                                                                    (STAR/)
(ROTH/)
(LARO/)
                                                                                                                                                                   (SHIM/)
(CRAB/)
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Sequence 20 BP; 11 A; 7 C; 0 G; 2 T; 0 U; 0 Other;
                     Key
modified_base
                                 modified_base
                                         modified base
               Homo sapiens
                                                       08-APR-2004
                                                                       Gierse JK;
                  Synthetic
                                                                                     ischemia.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to aga4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the mucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal natisense condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisinflammatory, neuroprotective, notoropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 expression of disease or condition associated with mpGES-1 expression of disease or condition associated with mpGES-1 inflammation, Alzheimer's disease, arthritis, diabetes, cancer, isolaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                            immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; natiantiinflammatory; neuroprotective; natiantiinflammatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
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residues are 5-methylcytidines"
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/note= "2'-0-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mCGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; imCGES-1; mPGES-1 inhibitor; intidabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1366,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "2'-0-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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                                                                                                     ADM15179 standard; DNA; 20 BP.
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19 GTATGTGTGTGTGTGT
                                                                                                                                                                        01-JUL-2004 (first entry)
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                      RESULT 1706
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16; Conservative

Local Similarity

Query Match Matches

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0; Gaps

Indels

2;

0; Mismatches

0.4%; Score 14.8; DB 1; Length 20; 38.9%; Pred. No. 1.7e+03;

88.9%;

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antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, antiarthritic, vasotropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nautionflammatory; neuroprotective; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1728.
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                                                                                                                                                                           Ouery Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                             2; Indels
                                                                                                                                              Sequence 20 BP; 11 A; 7 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                             0; Mismatches
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                          2315 GTCTGTGTGTGTGTGT 2332
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
                                                                                                                                                                                                                                                           20 GTATGTGTGTGTGTATGT 3
                                                                                                                                                                                                                                                                                                                                                    ADM15541 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                             16; Conservative
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modified_base
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                                                                                                                                                                                                             Matches
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New antisense compound, having a sequence targeted to a nucleic acid

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mEGES-1). The human meGES-1 gene is located on chromosome 9, more specifically too Gq44.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding meGES-1, which specifically hybridise with the nucleic acid meGES-1 and inhibits its expression; (2) a method of inhibiting the expression of meGES-1 in cells or tissues; and (3) a method of treating an animal condition associated with meGES-1. MPGES-1 chimeric antidabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound con be used as mPGES-1 inhibitors and in gene therapy. The antisense compound con the used as mPGES-1 inhibitors and in gene therapy. The antisense compound contion associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation alsorder.
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encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 10 A; 7 C; 0 G; 3 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "2'-0-methocyethyls"
16. .20
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                                                 Claim 4; SEQ ID NO 1728; 132pp; English.
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/mod.base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 88.9
Matches 16; Conservative
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                                         ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM14919;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1708
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/mcd_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

Location/Qualifiers

/*tag=

modified_base

Synthetic.

/mod_base= OTHER /note= "2'-0-methocyethyls" 16. .20

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1..5 /*tag=

modified base

/mod_base= OTHER /note= "2'-O-methoxyethyls"

/*tag=

modified_base

25-SEP-2003; 2003WO-US030374. 25-SEP-2002; 2002US-0413549P.

WO2004028458-A2

08-APR-2004

(PHAA) PHARMACIA CORP

WPI; 2004-305094/28.

Gierse JK;

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antidiabetic, immunomodulator, cardiant, neuroprotective, antiantlammatory, neuroprotective, nootropic, antianthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGBS-1 e.g., inflammation, Altheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8.30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoding
                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 1106; 132pp; English.
                                                                                                                                                            25-SEP-2002; 2002US-0413549P
                                                                                                                 25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                          (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                      WPI; 2004-305094/28
                                                                      08-APR-2004
                                                                                                                                                                                                                                                        Gierse JK;
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New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or

Claim 4; SEQ ID NO 1039; 132pp; English

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically. The human mpGES-1 gene is located on chromosome 9, more specifically to mpGES-1. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and antisense compounds have cytostatic, antidiabetic, immunomodulatory and cardiant activities, and can ophthalmological, immunomodulatory and cardiavascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 6 A; 9 C; 4 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2361 GTGTGCCTGTGTGCGTGC 2378
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Best Local Similarity 88.9
Matches 16; Conservative
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a<sub>K</sub>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1039,
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18 Grégéérigiérérérée 1

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2361 GIGIGCCIGIGIGGGIGC

Best Local Similarity 88.9%; Matches 16; Conservative

Query Match

ADM14852 standard; DNA; 20 BP

RESULT 1709 ADM14852/c

(first entry)

01-JUL-2004

ADM14852;

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Gaps

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ADM15032 standard; DNA; 20 BP.

ADM15032

Homo sapiens

can

Sequence 20 BP; 7 A; 8 C; 5 G; 0 T; 0 U; 0 Other;

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WPI; 2004-305094/28.
                     WO2004028458-A2
01-JUL-2004
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiathritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "phosphorothioate linkages and all cytidine
                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       residues are 5-methylcytidines
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/note= "2'-0-methocyethyls"
16. 20
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/note= "2'-O-methoxyethyls"
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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08-APR-2004

25-SEP-2003; 2003WO-US030374. 25-SEP-2002; 2002US-0413549P.

(PHAA) PHARMACIA CORP

Gierse JK;

New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or ischemia.

Claim 4; SEQ ID NO 1219; 132pp; English.

antidiabetic, immunomodulator, cardiant, neuroprotective, antiantlammatory, neuroprotective, nootropic, antianthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated whith mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder. encoding The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention also describes: (1) antisense compounds having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,

encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or

New antisense compound, having

WPI; 2004-305094/28.

a sequence targeted to a nucleic acid

The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to

Claim 4; SEQ ID NO 1597; 132pp; English.

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                                                                                                                                                                                                                         microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1; inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immicrosomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; aftidiation; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; albabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                            Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1597.
                       ö
                                                                                                                                                                                                                  antisense oligonucleotide; phosphorothioate; human;
 Length 20;
                      Indels
Score 14.8; DB 1;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .5
/*tag= a
/mod_base= OTHER
/note= "2'-0-methocyethyls"
                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                            2322 rerererererecerere 2339
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                                                                   18 rerererererere 1
                                                                                                                           BP.
 0.4%;
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                                                                                                                           ADM15410 standard; DNA; 20
                                                                                                                                                                      (first entry)
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                       Conservative
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          Similarity
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                                                                                                                                                                                                                                                                                                                                                                   Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified base
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Query Match
Best Local Simil
Matches 16; C
                                                                                                                                                                                                                                                                                                                                  sapiens.
                                                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-APR-2004,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gierse JK;
                                                                                                                                                                                                                  chimeric;
                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                 ADM15410;
                                                                                                    RESULT 1711
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                                                                                                                 ADM15410,
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9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8.30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. WFGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antinflammatory, neuroprotective, nootropic, aphthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                  disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; UCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acute respiratory distress syndrome; pulmonary hypertension;
lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ve 0; Mismatches 2; Indels
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Shahabuddin S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 10 A; 7 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                        2336 TGTGTGTGTGTGTGCA 2353
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23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                                                                                                                                                                                 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human oligonucleotide #1396
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                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 88.9
Marches 16; Conservative
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SANDRASAGRA A.
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SHAHABUDDIN S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JS2004049022-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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(TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1712
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The invention relates to oligonuclectides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin [L1]-4 receptor, interleukin [L1] chosen from a gene encoding interleukin [L1]-4 receptor, interleukin [L1] chosen from a gene encoding interleukin [L1]-4 receptor, interleukin [L1] chosen from a gene encoding interleukin [L2]-4 receptor, interleukin [L1] con encoding arget(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The prevention and/or treatment of a respiratory or lung disease. The colligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, ppB4 B, ppB4 C, or ppB4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor(s), and/or asthma and/or lung allergies associated with cifinamation or an inflammation, allergy, asthma, impeded respiration, allergy, asthma, impeded respiration, cycle ithrosis (CP), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory disease (COPD), allergic rhinitis, acute respiratory disease syndrome, pulmonary hypertension, lung inflammation, represents an oligonucleotide of the bronch constriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 B; PDE8 C; PDE8 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CODP; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertenaton; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 10 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                  Claim 2; SEQ ID NO 1397; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3196 CCGGAGCTGGAGGATCCC 3213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-APR-2002; 2002WO-US013135. 23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUL-2003; 2003US-00627930.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human oligonucleotide #1588.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19 crecaecrecaecec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADO46222 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JS2004049022-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ното варіелв.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO46222;
                                                      asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADO46222
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(MILL/) (SHAH/) (LUHH/) (CONG/)

Nyce JW,

asthma.

(SAND/) (TANG/)

(AGUI/)

NYCE/)

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The invention relates to antisense oligonuclectides targeted to a nucleic acid molecule encoding fatty acid synthase, which inhibit the expression of fatty acid synthase. The antisense oligonuclectides are useful for increasing the metabolic rate, decreasing adiposity, decreasing serum leptin, decreasing serum cholesterol, decreasing blood glucose, decreasing serum insulin, decreasing serum lipids and treating an animal having a disease or condition associated with fatty acid synthase, such as breast, prostate, colon, endometrium, ovary and thyroid cancers. It may also be useful prophylactically, e.g., to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human fatty acid synthase antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compound of 8-80 nucleobases in length which inhibits the expression of fatty acid synthase, useful for treating disease or condition associated with fatty acid synthase such as breast and colon cancers.
                                                                                                              /mod_base= Other
/note= "Phosphorothioate backbone. All cytidines are 5-
methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; fatty acid synthase; antisense therapy; metabolic rate; adiposity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
                                                                                                                                                                                                                                /*tag= a
/mod_base= Other
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                     /mod_base= Other
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human fatty acid synthase, antisense oligonucleotide #114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; SEQ ID NO 21; 87pp; English
                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2231 TAGCAGCCCACCCTGCTG 2248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bhanot S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 reschecceaccarecre 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-OCT-2002; 2002US-00274085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2002; 2002US-00274085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADO21409 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                 16. .20
                                                                                                                                                                                                                                                                                                                                               '*tag≈
                                                                                                      *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Freier SM, Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-340035/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (FREI/) FREIER S M.
(DOBI/) DOBIE K W.
(BHAN/) BHANOT S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004077570-A1
                                                 Key
modified_base
                                                                                                                                                                                                          modified base
                                                                                                                                                                                                                                                                                                                   modified base
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 mucleotides of the 5'-end or 3'-end of a mucleic acid farger chosen from a gene encoding interleukin (IL) -4 receptor, interleukin (IL) -5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDB4 A, PDB4 B, PDB4 C or PDB4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the oligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, EDE4 A, PDB4 B, PDB4 C, or PDB4 D. The oligonucleotides are useful for reducing or inhibiting expression of a complication or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine and/or asthma and/or lung allergies associated with materion or an inflammatory disease. The respiratory or lung disease is associated with hyper-respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), hypertension, lung inflammation, allergy, asthma, impeded respiration, hypertension, lung inflammation, bonchitis, airway obstruction or the bronchitis, airway obstruction or the propertion or an inflammation, allergy as an oligonucleotide of the bronchitis and or the sequence represents an oligonucleotide of the bronchitis and or the propertion or the propertion or the propertion or the propertion or an inflammation, allergy and propertion or the propertion of the propertion or an inflammation, and propertion or an inflammation or an inflammatio
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                                                                                                                                                                                                                                                                                                                                                                     Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; fatty acid synthase; antisense therapy; metabolic rate; adiposity; serum leptin; serum cholesterol; blood glucose; serum insulin; serum lipid; breast cancer; prostate cancer; colon cancer; endometrium cancer; ovary cancer; thyroid cancer; infection; inflammation; tumour; probe; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                         Tang L, Aguilar D, Miller S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human fatty acid synthase, antisense oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 6 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; SEQ ID NO 1589; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1573 CAGGTGGCCCGGGGCATG 1590
                                                                                                                                                                                                                  Sandrasagra A, Tany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 cedendececededeare 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-2004 (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
                        SANDRASAGRA A.
                                                                                                      MILLER S.
SHAHABUDDIN S.
                                                                                                                                                                                                                                                                                                                   WPI; 2004-293804/27.
                                                 TANG L.
AGUILAR D.
NYCE J W
                                                                                                                                                                                                                                                                 Shahabuddin S,
                                                                                                                                                                                 CONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention.
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RESULT 1714 AD021297

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vivlemore401-10.rng

WO2004038416-A1.

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                                                                                                                                                                                                                                                                                                                                                                                                                                         New compound of 8-80 nucleobases in length which inhibits the expression of fatty acid synthase, useful for treating disease or condition associated with fatty acid synthase such as breast and colon cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
serum leptin; serum cholesterol; blood glucose; serum insulin;
serum lipid; breast cancer; prostate cancer; colon cancer;
endometrium cancer; ovary cancer; thyroid cancer; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 1.76
0, Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 16; SEQ ID NO 133; 87pp; English
                                                             inflammation; tumour; probe; human.
                                                                                                                                                                                                                                                                                                                                                                      Freier SM, Dobie KW, Bhanot S;
                                                                                                                                                                                                                  17-OCT-2002; 2002US-00274085.
                                                                                                                                                                                                                                                      17-OCT-2002; 2002US-00274085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 88.9
Les 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-340035/31.
                                                                                                                                                                                                                                                                                        (PREI/) FREIER S M.
(DOBI/) DOBIE K W.
(BHAN/) BHANOT S.
                                                                                                                                       US2004077570-A1.
                                                                                                   domo sapiens
                                                                                                                                                                           22-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Matches
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RESULT 1717
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                                                                                                                                                                                                                                                                                                                        GPR6; GPCR; G-protein coupled receptor 6; antianaemic; haemostatic; cytostatic; cardiovascular; cardiant; vasotropic; antiarrhythmic; antiarteriosclerotic; hypotensive; CNS; antiparkinsonian; nootropic; neuroprotective; cerebroprotective; neuroleptic; anticonvulsant; nephrotropic; gastrointestinal; anti-inflammacory; antiulcer; anti-HIV; antiasthmatic; antiallergic; immunosuppressive; thyromimetic; dermatological; gene therapy; protein therapy; probe; ss.
                                                                                                                                                                                                                                                                               Human GPR6 receptor cDNA detecting probe 1.
2231 TAGCAGCCCACCCTGCTG 2248
                                                                                                                                                    ADO44492 standard; cDNA; 20 BP.
                                        20 rescaecceaccarecre 3
                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                       29-JUL-2004
                                                                                                                                                                                               AD044492
                                                                                                       RESULT 1716
ADO44492/C
XX
AC ADO44499.
XX
DT 29-UUL-',
XX
DY 29-UUL-',
XX
KW GPR6; GJ
KW CYtOStar
KW neuropric
KW neuropric
KW antiastl
KW antiastl
KW dermato
XX
HOMO Saj
OS Synthet:
XX
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Homo sapiens

Synthetic

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The invention relates to screening for therapeutic agents and involves contacting a test compound with a G-protein coupled receptor 6 (GPR6) contacting a test compound with a G-protein coupled receptor 6 (GPR6) polypeptide, and detecting blinding of the test compound to GFR6 as a compound concentration (or in the absence of the test compound or at the presence of a compound known to be a GFR6 polypeptide activity at a certain test compound known to be a GFR6 polypeptide regulator. The therapeutic agents are useful in treating disease such as hematological disorders of the peripheral and central nervous system, urological disorders of the peripheral and central nervous system, urological disorders of the peripheral and central nervous composition for treating disease such as hematological disorders, leukopenia, leukaemia, and lymphomas), cardiovascular disease (e.g., heart failure, myocardial infarction, ischemia, arkiythmias, atherosclerosis, and hypertensive vascular disease), disorders of the peripheral and central nervous system (e.g., Parkinson's disease, compaction, lupus nephritis, glomerulopathies, nephritis, computed is allored in disease, stroke, Alzheimer's disease, eleg., renal central nervous system (e.g., Parkinson's disease, contropical disorders (e.g., dysphagia, Barrett's metaplasia, creation conjunctivitis, and erectile dysfunction), carcerating, islet cropluctivitis, atrophy of gastric glands, gastric ulcers, chronic gastroenterological disorders (e.g., dysphagia, Barrett's metaplasia, chromic stress gastritis, atrophy of gastric glands, gastric disease, and conjunctivitis, hereditary angloedema, Hashimoto's thyroiditis, systemic lupus erythematosus and scleroderma, and central for children conjunctivitis, hereditary angloedema, Hashimoto's children conjunctivitis, hereditary angloedema, Hashimoto's amammal they are also useful for the regulation of GRBC crivity in a mammal to the mammal having the disease. The GRBC activity the present
                                                                                                                                                                                                                                                                                                                     Screening for therapeutic agents, useful for treating e.g. hematological diseases, comprises contacting a test compound with a G-protein coupled receptor 6 polypeptide and detecting binding of the test compound to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence represents a probe hybridising to the human GPR6 receptor cDNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; SEQ ID NO 4; 128pp; English
                                                                                                                                                                                                                                Golz S, Brueggemeier U, Summer H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2015 ACCTGGACCGTGTCCTTA 2032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HP.
                                                                                         15-OCT-2003; 2003WO-EP011394
                                                                                                                                    24-OCT-2002; 2002EP-00023769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative
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LD ADN37165 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-AUG-2004 (first entry)
                                                                                                                                                                                  (FARB ) BAYER HEALTHCARE
                                                                                                                                                                                                                                                                            WPI; 2004-365568/34
                                         06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                             polypeptide
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BXXXXEXB
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The present invention relates to antisense compounds targeted to a mucleic acid encoding human Gankyrin. The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of Gankyrin. The antisense oligonucleotide comprises at least one modified. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-methoxyethyl (2'-MOE) sugar moiety, the antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methylcycosine. The antisense oligonucleotide further comprises at least one modified nucleobase, useful for the treatment of diseases such as
                                                                                                                                                                                                                                                                                           New compound targeted to a nucleic acid molecule encoding Gankyrin and inhibits the expression of Gankyrin, useful for modulating the expression of Gankyrin or for diagnosing or treating, e.g. hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hyperproliferative disorders, e.g. cancer. The present sequence represents a human Gankyrin DNA target sequence for an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 7 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            Example 15; SEQ ID NO 105; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1. .20
/*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1447 GCGGCCAAGGGTAACCTG 1464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADO52173 standard; DNA; 20 BP.
                                                                                                                                                                21-NOV-2002; 2002US-00302027.
                                                                                                                                21-NOV-2002; 2002US-00302027.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCAGCCAAGGGTAACTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense technology; ss.
                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                             WPI; 2004-399721/37
                                                                                                                                                                                                                              Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 16; Conserv
                                                                 US2004102391-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide.
   cytostatic, ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-AUG-2004
                                                                                                 27-MAY-2004
                                                                                                                                                                                                                              Dean NM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADO52173;
                                                                                                                                                                                                                                                                                                                                              disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AD052173
   ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to antisense compounds targeted to a nucleic acid encoding human Gankyrin. The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of Gankyrin. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-0-methoxyethyl (2'-MOB) sugar moiety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methylcytosine. The antisense oligonucleotide further comprises such as hyperproliferative disorders, e.g. cancer. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compound targeted to a nucleic acid molecule encoding Gankyrin and inhibits the expression of Gankyrin, useful for modulating the expression of Gankyrin or for diagnosing or treating, e.g. hyperproliferative
                Antisense therapy; human; Gankyrin; hyperproliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense therapy; human; Gankyrin; hyperproliferative disorder; cancer;
                                                                                                                                                           /note= "This oligonucleotide has a phosphorothioate backbone and 2'-methyoxyethyl (2'-MOS) wings at the 5' and 3' ends, which are 5 mucleotides in length at each end. All cytidine residues are 5-methylcytidines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; rive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 6 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 45; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Gankyrin DNA target sequence #18.
                                                                                                Location/Qualifiers
                                 cytostatic; phosphorothioate; ss
                                                                                                                                               /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1447 GCGGCCAAGGGTAACCTG 1464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                          21-NOV-2002; 2002US-00302027
                                                                                                                                                                                                                                                                                                                                         21-NOV-2002; 2002US-00302027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 GCAGCCAAGGGTAACTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADN37225 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16; Conservative
                                                                                                                            *tag=
                                                                                                               .20
                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-399721/37
                                                                                                                                                                                                                                                                                                                                                                                                         Dean NM, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention
                                                                                                                                                                                                                                            US2004102391-A1.
                                                                                              Key
modified_base
                                                                Homo sapiens
                                                                                                                                                                                                                                                                            27-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADN37225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1718
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Gaps

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/note= "OTHER= Phosphorothioate backbone. All cytidines
47.
Human inhibitor of apoptosis-like antisense oligonucleotide seqid
                                                                                   cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; IAP-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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modified base
                                 modified_base
                                                                                                                                                                  29-APR-2004
 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADP76354;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADP76354
 8XELTTTTTTTTX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        δ
                                                                                                                                                                                                                                                                                                                                                                   The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding inhibitors of apoptosis (IAP) like, where the compound specifically hybridises with the nucleic acid molecule encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the expression of IAP-like and calls or tissues; screening for a modulator of IAP-like; a diagnostic method for identifying a disease state comprising in IAP-like; a call of IAP-like in a sample using at least one of the primers the presence of IAP-like in a sample using at least one of the primers of sequences comprising SEQ ID NO. 5 or 6, or the probect of comprising SEQ ID NO. 7; a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with IAP-like. The compound is useful for modulating the expression of IAP-like.

It is also useful for diagnosting or treating diseases associated with Corporation of IAP-like, e.g. a hyperproliferative disease. This sequence represents a human inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                 New compound targeted to a nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like and inhibits expression of IAP-like, useful for modulating the expression of IAP-like or for treating, e.g. hyperproliferative disorder.
                                 /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                           /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
reperfusion; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric phosphorothioate oligonucleotide #507
                                                                                                                                                                                                                                                                                                                                                 Example 14; SEQ ID NO 47; 58pp; English.
are 5-methylcytidines"
                                                                                base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3198 GGAGCTGGAGGATCCCCT 3215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADP76708 standard; DNA; 20 BP
                                                                                                                                                                                         22-NOV-2002; 2002US-00303325.
                                                                                                                                                                  22-NOV-2002; 2002US-00303325.

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                         ๙
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 88.9
Les 16; Conservative
                                                       15. .20
         /*tag=
                                                                      /*tag=
                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                       Bennett CF, Dobie KW;
                                                                                                                                                                                                                                                              WPI; 2004-399725/37.
                                                                                                                    US2004102395-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide
            modified base
                                                        modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP76708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1720
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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonuclectides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition. They are also useful in research and disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence respressors a chimeric phosphorothicate oligonuclectide with 2' MOE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase
(GFAI), for treating diabetes, a cardiovascular or neurologic disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 6 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric phosphorothioate oligonucleotide #153
                                                              /*tag= a
/mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                               /*tag= b
/mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 507; 175pp; English.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1473 TCTGCGGCGCGCGCC 1490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-OCT-2003; 2003WO-US03332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-OCT-2002; 2002US-0419268P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                17. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Broschat KO, Crosby SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-348453/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reperfusion; ss.
                                                                                                                                                                                                                                                                                                                                    WO2004035763-A2
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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MOE wings and a deoxy gap, these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/repertusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence 20 BP; 1 A; 6 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric phosphorothicate oligonucleotide #508
                                                                              /*tag= b
/mod_base= other
/note= "2-methoxyethyl wing"
                    /mod_base= other
/note= "2-methoxyethyl wing"
17. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 828; 175pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1473 TCTGCGGGCGCGCGCC 1490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADP76709 standard; DNA; 20 BP
                                                                                                                                                                                                                                                        02-OCT-2003; 2003WO-US03332
                                                                                                                                                                                                                                                                                                  17-OCT-2002; 2002US-0419268P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%;
Best Local Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 rerecesecresessece
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                       Broschat KO, Crosby SD;
                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-348453/32.
                                                                                                                                                                       WO2004035763-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reperfusion; ss.
                                                              modified base
                                                                                                                                                                                                               29-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP76709;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
  요
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the phresent sequence represents a chimeric phosphorothicate oligonucleotide with 2' MOE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New compounds, particularly antisense oligonucleotides targeted to a nuclaic acid encoding glutamine-fructose-6-phosphate amidotransferase (GRAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 1; Length 20;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 4 A; 2 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric phosphorothioate oligonucleotide #828
                                                                                  'note= "2-methoxyethyl wing'
                                                                                                                                                                     'note= "2-methoxyethyl wing'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 153; 175pp; English.
Location/Qualifiers
                                                              /mod base= other
                                                                                                                                               /mod_base= other
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GFAT; Antidiabetic; Cardiant;
                                                                                                                                                                                                                                                                                                                                             17-OCT-2002; 2002US-0419268P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%;
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Les 16; Conservative
                                                                                                    17. .20
                                                                                                                                                                                                                                                                                                                                                                                                                             Crosby SD;
                        ...4
*tag≃
                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                                                                                  (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-348453/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                reperfusion; ss.
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Key
modified_base
                                                                                                    modified base
                                                                                                                                                                                                                                                                                                                                                                                                                               Broschat KO,
                                                                                                                                                                                                                                                        29-APR-2004
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Gaps

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Query Match

Matches

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Location/Qualifiers 1. .4 /*tag= a /mod_base= other

Key modified_base

Location/Qualifiers 1. .4

Key modified_base

Synthetic

ADP77029;

RESULT 1722

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Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-MAY-2004
                                                            29-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1725
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      셤
                                                                                                                                                                                                                                                                        The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are artisense oligonuclectides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MCE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                     New compounds, particularly antisense oligonucleotides targeted to a
nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase
(GFAI), for treating diabetes, a cardiovascular or neurologic disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GFAT; Antidiabetic; Cardiant;
Glutamine-fructose-6-phosphate amidotransferase; diabetes; ischemia;
reperfusion; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 6 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric phosphorothioate oligonucleotide #285.
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/*tag= a
/mod_base= other
17..20
/note= "2-methoxyethyl wing"
                             /mod_base= other
/note= "2-methoxyethyl wing"
                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 508; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1473 TCTGCGGCGCGCGCC 1490
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP76486 standard; DNA; 20 BP
                                                                                                   02-OCT-2003; 2003WO-US03332
                                                                                                                      17-OCT-2002; 2002US-0419268P
                                                                                                                                                                                                                                     ischemia/reperfusion injury.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
          17. .20
/*tag= k
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                                                                                                                                                               Crosby SD;
                                                                                                                                          (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                   WPI; 2004-348453/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                            WO2004035763-A2
          modified_base
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                                                                                                                                                               Broschat KO,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    16;
                                                                               29-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADP76486;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonuclectides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MOE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
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inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 4 A; 2 C; 6 G; 8 T; 0 U; 0 Other;
/mod_base= other
/note= "2-methoxyethyl wing"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 285; 175pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3237 TAGTIGGAGGIGATICCA 3254
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                                                                                                                                                                                                                                                                                                                                      17-OCT-2002; 2002US-0419268P.
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20-DEC-2002; 2002US-00325899
                                                                                                                                                                                                                                                             02-OCT-2003; 2003WO-US03332
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Best Local Similarity 88.2
Best Local Similarity
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Broschat KO, Crosby SD;
                                                                                                                                                                                                                                                                                                                                                                                                       (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-348453/32.
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Query Match
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                                                                                                                                                                    The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an enotransplant rejection or mechanical organ replacement rejection, in an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other discorders.
                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                  Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human death-associated protein kinase 1 gene inhibitory oligo ISIS233836.
                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "phosphorothioate backbone, all C bases are 5-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; death-associated protein kinase 1; gene expression; diagnosis; dysregulation; cellular apoptosis.
                         Morris M;
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                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1; Length 20; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                               2; Indels
                         Prentice J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "2'-0-methoxyethyl nucleobase"
16. .20
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/mod_base= OTHER
/note= "2'-O-methoxyethyl nucleobase"
                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 2 C; 10 G; 4 T; 0 U; 0 Other;
                         Ľy N,
                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                               Claim 58; SEQ ID NO 795; 1762pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   methylcytidine bases"
                        Woodward R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
(EXPR-) EXPRESSION DIAGNOSTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                     853 GAGGAGGAGCTGGTGGAG 870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                   0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAGGTGGAGCTGGTGCAG
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                        Fry K,
                                                            WPI; 2004-400724/37.
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                        Wohlgemuth J,
Rosenberg S;
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modified_base
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                                                                                                                           genes.
                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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New compound targeted to a nucleic acid encoding death-associated protein kinase 1, useful for modulating death-associated protein kinase 1 expression, or treating diseases associated with expression of death-associated protein kinase 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a compound 8-80 nucleobases in length targeted where the compound specifically hybridizes with the nucleic acid molecule encoding death-associated protein kinase 1, where the compound specifically hybridizes with the nucleic acid molecule encoding death-associated protein kinase 1 and inhibits the expression of death-associated protein kinase 1. The compound is useful for the modulation of death-associated protein kinase 1 expression and for diagnosis and treatment of diseases associated with expression of death-associated protein kinase 1 expression. The disease or condition is dysregulation of cellular apoptosis. The compound is also useful in relationships that exist between death-associated protein kinase 1 and disease state, phenotype, or condition. This sequence represents an inhibitory oligonucleotide of the invention which is targeted to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human death-associated protein kinase 1 gene inhibitory oligo ISIS233836.
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methylcytidine bases"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.7e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human death-associated protein kinase 1 gene (ADO59470)
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/mod_base= OTHER
/note= "2'-O-methoxyethyl nucleobase"
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/note= "2'-0-methoxyethyl nucleobase"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 9 A; 2 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 25; SEQ ID NO 43; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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/mod_base= OTHER
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                                                                    22-NOV-2002; 2002US-00303588
21-NOV-2003; 2003WO-US037445
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATATATATATACATAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16; Conservative
                                                                                                                                         PHARM INC
                                                                                                                                                                                                                                                                           WPI; 2004-441167/41.
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modified_base
                                                                                                                                     SISI (-SISI)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-AUG-2004
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09-DEC-2002; 2002US-00316243.
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                                                                                   US2004110147-A1
             modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human BAF53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-AUG-2004
                                                                                                            10-JUN-2004.
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                                                                                                                                                                                                                           Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1729
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                                                                                                                                                                                                                                                                                                        The invention relates to a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding death-associated protein kinase 1, where the compound appecifically hybridizes with the nucleic acid molecule encoding death-associated protein kinase 1 and inhibits the expression of death-associated protein kinase 1. The compound is useful for the modulation of death-associated protein kinase 1. The compound is useful for the diagnosis and treatment of diseases associated with expression of death-associated protein kinase 1 expression. The disease or condition is dysregulation of cellular apoptosis. The compound is also useful in research and diagnostics, and for drug discovery to elucidate

This sequence represents and disease state, phenotype, or condition. This sequence represents and inhibitory oligonucleotide of the invention which is targeted to the human death-associated protein kinase 1 gene (ADO59470).
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                                                                                                                                                                                                            New compound targeted to a nucleic acid encoding death-associated protein kinase 1, useful for modulating death-associated protein kinase 1 expression, or treating diseases associated with expression of death-associated protein kinase 1.
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/note= "Phosphorothioate backbone and all cytidines are 5
-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; 88; antisense; chromatin; BAF complex; BRG1/brm-associated factor; BAF53; BRG1-associated factor 53kDa; cancer; tumour; actin-related protein; hyperproliferative disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 9 A; 2 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human BAF53 antisense oligonucleotide ISIS280402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                   Claim 25; SEQ ID NO 43; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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/mod base= OTHER
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                                                                    21-NOV-2003; 2003WO-US037445
                                                                                               22-NOV-2002; 2002US-00303588
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/*tag=
                                                                                                                          (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                   WPI; 2004-441167/41
                WO2004048531-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Key
modified base
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                                          10-JUN-2004
                                                                                                                                                        Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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The invention relates to a compound, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding BAF53 (a member of the BAF complex (BRG1/brm-associated factor 53kDa which is an actin-related protein), specifically hybridises with the nucleic acid encoding BAF53 comprising 28001-bp sequence (derived from human chromosome 3) and inhibits expression of BAF53, i.e. an antisense oligonucleoride. Also included are inhibiting the expression of BAF53 in cells or tissues, screening for a modulator of BAF53, a diagnostic method for identifying a disease state, a kit or assay device comprising the compound and treating an animal having a disease or condition associated with BAF53. The oligonucleotide compound is useful for preparing a timonum. The present sequence is an antisense oligonucleotide targeting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ds; antisense; chromatin; BAF complex; BRG1/brm-associated factor; BAF53; BRG1-associated factor 53kDa; cancer; tumour; actin-related protein; hyperproliferative disorder; chromosome 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligonucleotide compound that inhibits expression of BAF53, useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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'note= "2'-methoxyethyl residue"
                                                                                                 /mod_base= OTHER
/note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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Best Local Similarity 88.9%;
Matches 16; Conservative
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                     /*tag= c
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The invention relates to a microwell array chip for detecting a single antigen-specific lymphocyte. Bach microwell of the chip has the shape and size to accommodate one lymphocyte only. The lymphocyte may be a B or a T lymphocyte. The invention also relates to methods for detecting, isolating and selecting an antigen-specific receptor (e.g., an method of coning a gene encoding an antigen-specific receptor (e.g., an expect immunoglobulin or a T-cell receptor) from an antigen-specific lymphocyte wia reverse transcription-PCR (RT-PCR); a method of manufacturing or a monoclonal antibody using an antigen-specific immunoglobulin gene cloned using the cloning method of the invention; and a method of manufacturing gene therapy material using an antigen-specific T-cell receptor gene cloned using the cloning method. The method of the invention is useful for the efficient detection of a single antigen-specific lymphocyte, genes from which may be isolated and cloned for use in various communication and generally methods. Sequences App21186-ADP21233 response the PCR primers used to clone the heavy and light chain variable regions.
                                                                                                         Microwell array chip for detecting antigen specific lymphocyte, has shape and dimension to store lymphocyte in microwell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide compound that inhibits expression of alpha-methylacyl -CoA racemase, useful for preparing a composition for treating a condition involving defects in fatty acid metabolism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         alpha-methylacyl-CoA racemase; AMACR; fatty acid metabolism;
gene therapy; ds; human; antisense therapy target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 14.8; DB 1; Length 20; 80.0%; Pred. No. 1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 1 A; 7 C; 5 G; 6 T; 0 U; 1 Other
                  Suzuki M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                                                                                                                                                        Example 5; SEQ ID NO 4; 92pp; Japanese
                  Tamiya E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       932 TCATCCTGGTGGTGGCGCT 951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 TCCTCCTSCTGGTGGCAGCT 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-DEC-2003; 2003WO-US039230.
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nes 16; Conservative
                Muraguchi A, Kishi H,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP56824;
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                                                                                                                                                                                                                                                                                                                             The invention relates to a compound, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding BAF53 (a member of the BAF complex (BRG1/Dram-associated factor), BRG1-associated factor 53Mpa which is an actin-related protein), specifically hybridises with the nucleic acid encoding BAF53 comprising 28001-bp sequence (derived from human chromosome 3) and inhibits expression of BAF53, i.e. an antisense oligonucleotide. Also included are inhibiting the expression of BAF53 in cells or tissues, screening for a modulator of BAF53, a diagnostic method for identifying a disease state, a kit or assay device comprising the compound and treating an animal having a disease or condition associated with BAF53. The oligonucleotide compound is useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer or a tumour. The BAF53 gene is located on chromosome 3. The present sequence of a number of the contract of the contra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               is a BAF53 genomic DNA target sequence for an antisense oligonucleotides.
                                                                                                                                                                     New oligonucleotide compound that inhibits expression of BAF53, useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Heavy chain variable region (VH) 5' PCR primer hVH4a.1, SEQ:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 0.4%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.7e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 1 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                        Example 15; SEQ ID NO 167; 72pp; English.
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                  09-DEC-2002; 2002US-00316243.
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29-NOV-2002; 2002JP-00346728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-2004 (first entry)
                                                               (ISIS-) ISIS PHARM INC
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                                                                                                                                                     WPI; 2004-440336/41
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TAMIYA E.
SUZUKI M.
                                                                                                           Jain
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                                                                                                           Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP21189;
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FRIEDMAN P N. GORDON J. HODGES R. C. KLASS M R. KRATOCHVIL J D. ROBERTS- RAPP L. RUSSELL J C. STROUPE S D.
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08-OCT-1997;
15-OCT-1999;
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                                                                                                   Query Match
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(KLAS/)
(KRAT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (COHE/)
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                                                                                                                                                                                                                                                       RESULT 1733
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide compound that inhibits expression of alpha-methylacyl -CoA racemase, useful for preparing a composition for treating a condition involving defects in fatty acid metabolism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
                            The invention relates to a novel compound having a sequence comprising 8-80 bp which is targeted to a nucleic acid encoding alpha-methylacyl-COA racemase (AMACR), specifically hybridises with the nucleic acid encoding alpha-methylacyl-COA racemase and inhibits the expression of alpha-methylacyl-COA racemase. The oligonucleotide compound of the invention may be useful for preparing a composition for treating a disease or condition involving defects in fatty acid metabolism, possibly via gene therapy. The current sequence is that of a human AMACR DNA of the invention which was targeted for antisense therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel compound having a sequence comprising 8-80 bp which is targeted to a nucleic acid encoding alpha-methylacyl-CoA racemase (AMACR), specifically hybridises with the nucleic acid encoding alpha-methylacyl-CoA racemase and inhibits the expression of alpha-methylacyl-CoA racemase. The oligonuclectide compound of the invention may be useful for preparing a composition for treating a disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "OTHER = Bases 1-5 and 16-20 are 2'-methoxyethyl (2'-MOE) bases, all cytidines are 5'-methylcytidines, phosphorothioate backbone throughout"
                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense 2'-MOE gapmer targeted to human AMACR RNA - SEQ ID 18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 alpha-methylacyl-CoA racemase; AMACR; fatty acid metabolism; gene therapy; antisense; 2'-methoxyethyl gapmer; 2'-MOE wing; phosphorothioate backbone; 8s; human.
                                                                                                                                                                                                                                    0.4%; Score 14.8; DB 1; Length 20; 88.9%; Pred. No. 1.78+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                      Sequence 20 BP; 3 A; 7 C; 8 G; 2 T; 0 U; 0 Other;
 Example 15; SEQ ID NO 95; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 15; SEQ ID NO 18; 157pp; English.
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                                                                                                                                                                                                                                                                                                           3659 CCTGCAGGCCATGCCTC 3676
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                                                                                                                                                                                                                                                  Local Similarity
les 16; Conserv
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Matches
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condition involving defects in fatty acid metabolism, possibly via gene therapy. The current sequence is that of a antisense 2'-methoxyethyl (2'-MOE) gapmer oligo of the invention which was targeted to human AMACR RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting a target PS112 polynucleotide, useful in diagnosing, staging, monitoring, prognosticating, preventing and treating prostate cancer, comprises contacting the test sample with PS112-specific polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic, gene therapy, PS112, recombinant expression system; PS112 epitope; prostate disease; tumours; metastasis; predisposition; prostate cancer; RT-PCR; reverse transcriptase PCR; primer; 85.
                                                                                                                                           Gaps
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                                                                                                      0.4%; Score 14.8; DB 1; Length 20;
88.9%; Pred. No. 1.7e+03;
ve 0; Mismatches 2; Indels
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Stroupe SD,
                                                                    Sequence 20 BP; 2 A; 8 C; 7 G; 3 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nan PN, Gordon J, Hodges SC,
Roberts-Rapp L, Russell JC,
                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                  Human PS112 RT-PCR primer seqid 21.
                                                                                                                                                                               3659 CCTGCAGGCCATGGCTC 3676
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                                                                                                                                                                                                                                                                                                          BP.
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97US-00946869.
99US-00418887.
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                                                                                                                         Best Local Similarity 88.9%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                          ADQ96633 standard; DNA; 20
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cc that includes an open reading frame derived from PS112 operably linked to a control sequence compatible with a desired host, where the nucleic acid sequence has at least 50% identity to a sequence of SEQ ID NOS: 1-10, or their fragments or complements; a cell transfected with the recombinant expression system or with a nucleic acid sequence encoding at least one PS112 epitope, where the nucleic acid sequence comprises SEQ ID NOS: 1-0, or their fragments or complements; a composition of matter comprising a PS112 polynucleotide or its fragment, where the polynucleotide has at composition of matter comprising a PS112 polynucleotide or its fragment, where the polynucleotide has at complements, or has at least 50% identity with fragments of a complements or bas at least 50% identity with SEQ ID NOS: 2-10, or their complements or has at least 50% identity with SEQ ID NOS: 9 or 10. The method is DNA having at least 50% identity with SEQ ID NOS: 9 or 10. The method is custing the presence of a target PS112 polynucleotide in a test sample. The methods, test kit, polynucleotides and polypeptides, and continued in detecting and treating prostate diseases, tumours or complements are useful in detecting, diagnoshing, staging, monitoring, prognosticating, preventing and treating prostate diseases, tumours or complements are verse transcriptase PCR primer used to isolate conditions of the prostate, e.g. prostate cancer. This colate conditions of the prostate, e.g. prostate cancer. This colate conditions of the prostate of the prostate cancer. The colate colate cancer transcriptase PCR primer used to isolate transcriptase prostate cancer.
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.; 0 Length 20; 2; Indels Sequence 20 BP; 0 A; 9 C; 0 G; 11 T; 0 U; 0 Other; Score 14.8; DB 1; Pred. No. 1.7e+03; 0; Mismatches 938 Cricificarcricarica 19 0.4%; 921 CTICTICCIGITCATCCT 16; Conservative Local Similarity N Query Match Matches g 8

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Gaps

AAQ25299 standard; DNA; 21 (revised) 25-MAR-2003 23-DEC-1992 AAQ25299;

BP.

ODN6 - control oligonucleotide. (first entry)

Transcription; inhibition; duplex; anti-sense therapy; RNA; polymerase; triple helix; triplex; ss

Synthetic.

WO9210590-A1

25-JUN-1992

91WO-US009321 10-DEC-1991;

90US-00625680 10-DEC-1990;

(GILE-) GILEAD SCI INC

Toole JJ;

WPI; 1992-234645/28.

Inhibiting transcription of duplex DNA in anti-sense therapy and diagnosis - by contacting the transcribed region of DNA with an oligomer to form triple helix. diagnosis

rable 1; Page 26; 45pp; English.

The oligomer is used as a control oligonucleotide for studying inhibition of transcription of duplex DNA. Similar oligomers are capable of binding to the transcribed region of the DNA so as to form a triple helix. This interaction is compared to the binding of the control oligonucleotide to

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give accurate results. The target region may lie within an exon or an intron and the oligomer forms a triple helix by explicting the GT motif (i.e. the oligomer is useful in antisense therapy. It can also be used (opt. in labelled form) diagnostically to detect target DNA or RNA by hybridisation. See also AAQ25290-300.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Heptad; plants; custom tailored storage proteins; in vivo; expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic polypeptide(s) contg. specified heptad units - expressed in vivo in plants to serve as custom-tailored storage proteins with specified aminoacid content.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          used to transform plants to increase the content of partic. amino acid
such as lysine or methionine in seeds or leaves. See also AAQ36810-28,
AAQ37265-301. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                        Sequence 21 BP; 0 A; 0 C; 11 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                      (Updated on 25-MAR-2003 to correct PN field.)
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(first entry)
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22-JUN-1993
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DNA sequencing; amplification; polymerase chain reaction; primer;
ligase chain reaction; thermal cycle labelling; PCR; ss.
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les 16; Conservative
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                                                                                                                                                                    Skowron PM,
                                                                                                                                                                                         WPI; 1996-251756/25.
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                                                                                                                      04-NOV-1994;
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                                                                                                 03-NOV-1995;
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                                                      WO9614405-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-NOV-1996
                                                                           17-MAY-1996
                                                                                                                                                                   Mueller RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .7-MAY-1996
                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT27276;
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1738
                                                                                                                                                                                                                                                                                                                                                                                                         Best Loca
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide SM90 (AAQ94988) and complementary sequence SM91 (AAQ94989) code for heptad peptide SSP10 (AAR78247). They were annealed and used in the construction a DNA fragment (see also AAQ94996) that was inserted into vector pSK6 (see also AAR78316). The DNA fragment codes for a synthetic storage protein (SSP) contg. multiple lyaine-rich heptad repeats (see AAR78253). This can be expressed in the seeds of transformed plants, e.g. soybean and corn, to increase lysine content
                                                                                                                                                                                                                                                                                                                                                                                                   New chimeric gene providing increased lysine content in plant seeds - contains di:hydro:di:picolinic acid synthase gene coupled to chloroplast transport sequence and seed specific promoter, also new plants of improved nutritional value.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA polymerase I; truncated; exonuclease free; holoenzyme; universal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                            Lysine, synthetic storage protein; SSP; vector; pSK6; dihydrodipicolinic acid synthase; corn; maize; Zea mays; soybean; Glycine max; transgenic plant; essential amino acid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer FTFLM for DNA polymerase I gene primer walking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 2 A; 9 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                      (DUPO ) DU PONT DE NEMOURS & CO E I.
                                                                                                                                                                                                                /standard_name= "SM
                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 8; Page 78; 180pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1353 GGAGATGATGAAGATGAT 1370
                                                                                                                                                                                                                                                                                                                                                              Keeler SJ, Rice JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                      AAQ94989 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                  94US-00261661.
                                                                                                                                                                                                                                                                                 94WO-US013190
                                                                                                                                                                                                                                                                                                      93US-00160117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 GGAGAAGATGAAGAAGAT
                                                                                       SSP10 Oligonucleotide SM 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT27275 standard; DNA; 21
                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 88.9
les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-215272/28
                                                                                                                                                                                                                                     W09515392-A1
                                                                                                                                                                                                                                                                                 21-NOV-1994;
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                                                                15-JUL-1996
                                                                                                                                                                                          misc_feature
                                                                                                                                                                                                                                                                                                                  17-JUN-1994;
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                                                                                                                                                                                                                                                           08-JUN-1995
                                                                                                                                                                                                                                                                                                                                                             Falco SC,
                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT27275;
                                            AAQ94989;
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RESULT 1736
AAQ94989/c
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ID AAT2
XX AAT2
AC AAT2
XX 29-N
XX DY 29-N
XX DY KW DNA
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polymerase coding sequence, e.g. an exonuclease-free fragment (Tfl exo-fragment). This recombinant polymerase is thermostable and can be used in applications such as DNA sequencing, polymerase chain reaction, (universal) thermal cycle labelling and ligase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT27263-88 were synthesised for primer walking to obtain the remainder of the sequence of the Thermus flavus DNA polymerase I (holoenzyme) gene. The Tfl DNA pol I sequence can be used to generate a truncated DNA
                                                                                                                                                                                                                              Biologically active fragments of Thermus flavus DNA polymerase - useful for DNA sequencing, polymerase chain reaction, thermal cycle labelling and ligase chain reaction, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA polymerase I; truncated; exonuclease free; holoenzyme; universal; DNA sequencing; amplification; polymerase chain reaction; primer; ligase chain reaction; thermal cycle labelling; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.8; DB 1; Length 21; 38.9%; Pred. No. 1.8e+03; Ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer RTFLN for DNA polymerase I gene primer walking.
                                                                               RF;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 7 A; 2 C; 10 G; 2 T; 0 U; 0 Other;
                                                                           Piehl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                               Swaminathan N,
(MOLE-) MOLECULAR BIOLOGY RESOURCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MOLE-) MOLECULAR BIOLOGY RESOURCES INC
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Sequence 21 BP; 4 A; 2 C; 11 G; 4 T; 0 U; 0 Other;

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AAT27263-88 were synthesised for primer walking to obtain the remainder of the sequence of the Thermus flavus DNA polymerase I (holoenzyme) gene. The Tfl DNA pol I sequence can be used to generate a truncated DNA polymerase coding sequence, e.g. an exonuclease-free fragment (Tfl exofragment). This recombinant polymerase is thermostable and can be used in applications such as DNA sequencing, polymerase chain reaction, (universal) thermal cycle labelling and ligase chain reaction,
           Biologically active fragments of Thermus flavus DNA polymerase - useful
                       DNA sequencing, polymerase chain reaction, thermal cycle labelling
                                                                                                                                                                                                                                                                                                                                                                                                                                 Ovarian; epithelial cell line; drug screening; cancer research; human;
                                                                                                                                                                                                      0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03;
                                                                                                                                                                               Sequence 21 BP; 2 A; 10 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                            Human genomic DNA exon 7 amplifying primer 2.
                                                                                                                                                                                                                              0; Mismatches
                                                          Example 5; Page 45; 139pp; English.
                                                                                                                                                                                                                                                     2701
                                                                                                                                                                                                                                                                                                                    tumour; genomic; PCR primer; ss.
                                    and ligase chain reaction, etc.
                                                                                                                                                                                                                                                                 2 TCCAGGCTTCCCCACTCC 19
                                                                                                                                                                                                                                                    2684 TCCAGGCTTTCCCACTTC
                                                                                                                                                                                                                             16; Conservative
                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                               RESULT 1739
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Gaps ö

2; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        normal somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometriosis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                              Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dypplastic lesion; benjan tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                             Gaps
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    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03;
                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 12 A; 2 C; 0 G; 7 T; 0 U; 0 Other;
0.4%; Score 14.8; DB 1;
ilarity 88.9%; Pred. No. 1.8e+03;
Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stanton VP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Fig 7; 605pp; English
                                                                                        2695 CCACTTCCCACCCTGCCC 2712
                                                                                                                              21 ccacrreccaccerecae 4
                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                             Human polymorphic region 289.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-US005419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-0041057P.
                                                                                                                                                                                                                                   AAZ26100 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-521232/44.
  Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                     30-NOV-1999
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                                                                                                                                                                                                                                                                            AAZ26100;
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                                                                                                                                                                                          RESULT 1740
                                                                                                                                                                                                                  AAZ26100,
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Best Local Similarity

This primer is used for the PCR amplification of the exon 7 of human epithelial Cell lines 1040-21G (ATCC CRL 11730), TOV-112D (ATCC CRL 11731), TOV-112D (ATCC CRL 11731), OV-90 (ATCC CRL 11732), TOV-112D (ATCC CRL 11731), OV-90 (ATCC CRL 11732) and NOV-31 (ATCC CRL 11733). These cell lines which are new function as models for studying the ovarian expithelium under normal and pathological conditions, to assess the expression of genes involved in tumour suppression or tumour promotion. The cell lines can be used to screen for drugs that influence matrix metalloprotease activity and to screen for compounds useful in human or veterinary medicine through their influence on tumourogenicity, invasive potential or metalloprotease activity

ovarian epithelial cell lines - useful for cancer research, drug

Disclosure; Col 19-20; 19pp; English.

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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present in a population and targets only one allelic form present in the normal somatic cells, and the first gene. The products and methods can be used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometriosis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AZZS812-Z26825 represent
                                                                                                                                                                                                                                                                                                                                                               Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; prevention metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human polymorphic sites described in the method of the invention
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  2; Indels
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stanton VP;
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                                                                                                                                                                                          BP.
                                                                                  18 rrraggaarrrrarrrr 1
                                                                                                                                                                                                                                                                                                                         Human polymorphic region 290.
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                                          3316 TTTAGGAGATTTATTTT
                                                                                                                                                                                          AAZ26101 standard, DNA; 21
                                                                                                                                                                                                                                                                               30-NOV-1999 (first entry)
  16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (VARI-) VARIAGENICS INC
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  Matches
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used for preventing the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor present in a population and targets only one allelic forms of the gene normal somatic cells, and the first gene. The products and methods can be normal somatic cells, and the first gene. The products and methods can be used in the diagnosals, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic lesions, benign tumours, endometriosis, polycystic kidney disease, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                     cell viability, loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atheroscierotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                      Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stanton VP;
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                                                                                                    BP
18 TTTAGGAATTTTATTTT 1
                                                                                                                                                                                                                                 Human polymorphic region 994.
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                                                                                                    AAZ26805 standard; DNA; 21
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Best Local Similarity 88.9
Matches 16; Conservative
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Gaps

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Gaps

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3316 TTTAGGAGATTTATTTT

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Genetic proximity, gene expression, cell characterisation, homeobox gene, genetic defect, reverse transcriptase polymerase chain reaction, RT-PCR, kinase gene, protein phosphatase, P450, steroid receptor, cadherin,

Homo sapiens. WO9934016-A2

primer; ss Synthetic

PTK 11 gene specific primer.

(first entry)

11-0CT-1999

AAZ18112;

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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is transformed. They can be used for determining the effect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired containing cells capable of expressing an homeobox related desired (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZIN803-z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The
                                                                                                          Genetic proximity, gene expression, cell characterisation, homeobox gene; genetic defect, reverse transcriptase polymerase chain reaction, RT-PCR, kinase gene, protein phosphatase, P450, steroid receptor, cadherin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      family can be selected from a set of homeobox genes, kinase genes,
                                                                                                                                                                                                                                                                                                                                                                                                           Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein phosphatase genes, P450 enzyme genes, steroid receptor superfamily genes or cadherin superfamily genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 43; 102pp; English.
           AAZ18122 standard; DNA; 21 BP
                                                                                                                                                                                                                                                            98WO-IL000625
                                                                                                                                                                                                                                                                                   97IL-00122793
98IL-00126627
                                                                                    PTK 16 gene specific primer
                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-419113/35.
                                                                                                                                                                                                                                                                                                                       (GENE-) GENENA LTD
                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAY14657
                                                                                                                                                                                     Homo sapiens.
                                                             11-OCT-1999
                                                                                                                                                                                                           WO9934016-A2
                                                                                                                                                                                                                                                            28-DEC-1998;
                                                                                                                                                                                                                                                                                 29-DEC-1997;
16-OCT-1998;
                                                                                                                                                                                                                                   08-JUL-1999.
                                                                                                                                                 primer; ss.
                                                                                                                                                                       Synthetic.
                                     AAZ18122;
                                                                                                                                                                                                                                                                                                                                               Vider B;
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Identifying and characterizing cells by comparing the pattern of gene

expression in a selected gene family.

WPI; 1999-419113/35. P-PSDB; AAY14647.

97IL-00122793. 98IL-00126627. 98WO-IL000625

28-DEC-1998; 29-DEC-1997; 16-OCT-1998;

08-JUL-1999,

(GENE-) GENENA LID.

Vider B;

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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell compises: (a) obtaining the first cell and the second cell compises: (a) obtaining the first cell and the second cell the pattern cell; (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a compositive index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its characterising cells, e.g. for determining the origin of a cell, its contained. They can also be used for determining the cit an individual, e.g. a fetus. They can also be used for determining the ceffect of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired property. The method uses reverse transcriptase polymerase chain reaction (RP-PCR) for determining the pattern of gene expression in a selected on the RP-PCR reactions to determine the pattern of gene expression. The can be used from a set of homeobox genes, kinase genes, protein phosphatase genes, P450 enzyme genes, steroid receptor
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88.9%; Pred. No. 1.8e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 9 A; 7 C; 3 G; 2 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    superfamily genes or cadherin superfamily genes
Claim 4; Page 42; 102pp; English.
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nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ18114;
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Gaps ö

2; Indels

0; Mismatches

88.98;

Best Local Similarity 88.9 Matches 16, Conservative

Query Match

1801 GACGTCTGGTCCTTTGGG 1818

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18 caccicracitoricai

AAZ18112 standard; DNA; 21 BP.

AAZ18112/ ID AAZ1 XX

RESULT 1744

0.4%; Score 14.8; DB 1; Length 21; 18.9%; Pred. No. 1.8e+03;

Genetic proximity, gene expression, cell characterisation, homeobox gene, genetic defect, reverse transcriptase polymerase chain reaction, RT-PCR; kinase gene, protein phosphatase, P450, steroid receptor, cadherin,

98WO-IL000625 97IL-00122793. 98IL-00126627.

28-DEC-1998; 29-DEC-1997; 16-OCT-1998;

08-JUL-1999

Homo sapiens. WO9934016-A2.

primer; 88. Synthetic

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The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and a second cell comprises: (a) obtaining the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in the first cell and the second cell; (b) determining in a selected gene family; and (c) calculating a proximity index using a specified formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its genetic status, whether it carries a genetic defect, or whether it is cransformed. They can be used for determining the origin of a selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired containing the method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZINS03-ZIS332 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The
The invention provides a new method for identifying and characterising cells. The method for determining the genetic proximity of a first cell and as ascond cell comprises: (a) obtaining the first cell and the second cell comprises: (a) obtaining the first cell and the second cell comprises: (b) determining in the first cell and the second cell the pattern of expression of genes in a selected gene family; and (c) calculating a profifted formula. The methods can be used for characterising cells, e.g. for determining the origin of a cell, its centransformed. They can be used for detecting a selected genetic defect in an individual, e.g. a ferus. They can also be used for determining the obtaining the captes of the selected treatment on a test cell. They can also be used for obtaining cells capable of expressing an homeobox related desired compression in a selected property. The method uses reverse transcriptase polymerase chain reaction (RT-PCR) for determining the pattern of gene expression in a selected gene family. Sequences AAZ1780-Z18342 represent primers that can be used in the RT-PCR reactions to determine the pattern of gene expression. The gene family can be selected from a set of homeobox genes, kinase genes, conversally can be selected from a set of homeobox genes, kinase genes, increasing the pattern of genes, steroid receptor
                                                    Genetic proximity; gene expression; cell characterisation; homeobox gene; gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 6 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        superfamily genes or cadherin superfamily genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 42; 102pp; English.
                                                                                                                                                                                                                                                                                            98WO-IL000625.
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                                                                                                                                                                                                                                                                                                                                                   98IL-00126627
                  PTK 12 gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-419113/35.
                                                                                                                                                                                                                                                                                                                                                                                          (GENE-) GENENA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P-PSDB; AAY14649
                                                                                                                                                                                                           WO9934016-A2.
                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                          28-DEC-1998;
                                                                                                                                                                                                                                                                                                                                 29-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                     16-OCT-1998;
                                                                                                                                                                                                                                                  08-JUL-1999
                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                 Vider B;
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Identifying and characterizing cells by comparing the pattern of gene expression in a selected gene family.

WPI; 1999-419113/35.

Vider B;

P-PSDB; AAY14631.

(GENE-) GENENA LTD.

Claim 4; Page 42; 102pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                           Genetic proximity, gene expression; cell characterisation; homeobox gene; genetic defect; reverse transcriptase polymerase chain reaction; RT-PCR; kinase gene; protein phosphatase; P450; steroid receptor; cadherin;
                                                                                                                           Gaps
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protein phosphatase genes, P450 enzyme genes, steroid receptor
                                                                                    Score 14.8; DB 1; Length 21;
Pred. No. 1.8e+03;
0; Mismatches 2; Indels
                                                  Sequence 21 BP; 9 A; 7 C; 3 G; 2 T; 0 U; 0 Other;
                 superfamily genes or cadherin superfamily genes
                                                                                                                           0; Mismatches
                                                                                                                                                               1800 TGACGTCTGGTCTTTGG 1817
                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                                                              BP.
                                                                                      Query Match 0.4%;
Best Local Similarity 88.9%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                            PTK 8 gene specific primer.
                                                                                                                                                                                                                                                                             AAZ18106/c
ID AAZ18106 standard; DNA; 21
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Gaps

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1801 GACGICTGGTCCTTTGGG 1818

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Local Similarity 88.9 ses 16; Conservative

Best Loca Matches

Query Match

caccingrecriccinces 1

18

AAZ18096 standard; DNA; 21 BP

PTK 3 gene specific primer

(first entry)

11-OCT-1999

AAZ18096;

RESULT 1746
AAZ18096/C
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AAZ18096
AC
AC
AC
AZ18096
XX
AZ18096
XX
YX
XX
XX
XX
XX
XX

vivlemore401-10.rng

Vider B;

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A method has been developed of determining the Human Leukocyte Antigen class II (HLA Class II) group type of a subject. The method comprises:

(1) amplifying a target DNA sample from a subject using a particular HLA group-specific primer pair (sequence specific primer PCR - SSP-PCR); and (ii) determining whether a nucleic acid product is produced, therefore identifying the group type. AAX00303 to AAX0030s represent specifically claimed oligonucleotide primer for use in the above method. These coligonucleotide primer for use in the above method. These coligonucleotides are useful for determining the HLA Class II type of a patient sample, by identifying the specific alleles present and determining the group specificity of alleles. Steps (i) and (ii) in the method are diagnostically useful for histocompatibility analysis to see if donor and recipient groups match. The new sequences are useful for providing an insight into the genetic relationship between different corporation and recipient for group match can be grounded and alleles as few exon sequences offer conserved methods using exon based primers is more efficient than prior art methods using exon based primers, as few exon sequences offer conserved primer binding sites, resulting in a limited number of primer pairs and insufficient specificity for alleles, as allelic variations exist between the primer sites. The SSP-PCK method allows separation of haplotypes in the primer sites. The SSP-PCK method allows separation of cis-trans linkages of the patient samples, allowing resolution of cis-trans linkages of
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determining the Human Leukocyte Antigen Class II type Histocompatibility antigens - by using new intron-specific oligonucleotide primers for sequence specific primer PCR and sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary bypertension; pulmonary vasoconstriction; emphysema; ohronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            heterozygous sequencing results which cannot be achieved with other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1; Length 21;
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 7 A; 8 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Adenosine A2b receptor antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Fig 3; 36pp; English.
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                                                                                                                                  98EP-00111696
                                                                                                                                                                                                   97EP-00110438
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Local Similarity 88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-083585/08.
                                                                                                                                                                                                                                                                (BIOT-) BIOTEST AG.
                                                                                                                                                                                                   26-JUN-1997;
                                                                                                                                  25-JUN-1998;
EP892069-A2.
                                                                 20-JAN-1999
                                                                                                                                                                                                                                                                                                                                 Blasczyk R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human leukocyte antigen class II type; HLA class II type; histocompatibility locus antigen class II; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           superfamily genes or cadherin superfamily genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX00349 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                               98WO-IL000625
                                                                                                                                                                                                                                                                                                                                                             97IL-00122793.
98IL-00126627.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-419113/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENE-) GENENA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAY14641.
                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                            28-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                         29-DEC-1997;
                                                                                                                                                             WO9934016-A2
                                                                                                                                                                                                                                                                                                                                                                                               16-OCT-1998;
                                                                                                                                                                                                                              08-JUL-1999
primer; 88
                                                             Synthetic
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Gaps

; 0

2; Indels

Query Match

Matches

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Homo sapiens.

Synthetic.

AAX00349;

RESULT 1748
AAX 00349/c
XX
AC AAX00345
AC AAX 00345
DT 23-APR-1
XX
XX
Human 1e
XX

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Antisense oligonucleotides (ASO) for inhibiting a tumour necrosis factoralpha) gene in an animal, preferably a human, comprise 12-50 nucleotides, 90% of which are complementary to a region of mRNA containing a GGGA sequence motif. The ASO is used to inhibit expression of a gene in an animal and for treating the animal when afflicted with a disease or disorder characterised by the presence of an mRNA from a gene containing a GGGA motif. The ASO are specifically targeted to a GGGA sequence motif found in mRNA from a gene. A study of known ASO has shown that at least half of the most efficacious ASO's contain one or more TCCC against This ASO is designated TUU-0656 and is based on and targetted against the rat TNF-alpha sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human ABC1 cholesterol transporter; chromosome 9q31; ATP-binding cassette; HDL deficiency disorder; hlgh density lipoprotein; Tangier disease; TD; familial HDL deficiency; FHA; polymorphism; cardiovascular disease; coronary artery disease; coronary restenosis; cerebrovascular disease; coronary artery disease; coronary restenosis; cerebrovascular disease; peripheral vascular disease; Albachimer's disease; X-linked adrenoleukodystrophy; cancer; gene therapy; genetic diagnosis; prophylaxis; drug screening; transgenic animal; ds.
                                                                                                                                                                                                                                                                                                          Generation of antisense oligonucleotides - by specifically targeting a GGGA motif found in mRNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1; Length 21; Pred. No. 1.88+03; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ABC1 gene promoter polymorphic site, SEQ ID NO:204.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 0 A; 8 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 40; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1806 CTGGTCCTTTGGGGTCCT 1823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 credriccriredratici 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0124702P.
99US-0138048P.
99US-0139600P.
99US-0151977P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC69305 standard; DNA; 21 BP.
                                                                                                                                                                (UYJE-) UNIV JEFFERSON THOMAS
                                                                   98WO-US013711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAR-2000; 2000WO-IB000532
                                                                                                                97US-0051705P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JAN-2001 (first entry)
                                                                                                                                                                                                                                                               WPI; 1999-105767/09.
                                                                                                                                                                                                                Israel Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200055318-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                      02-JUL-1998;
                                                                                                                03-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAR-1999;
08-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUN-1999
                      14-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC69305;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1751
                                                                                                                                                                                                                  Τ̈.
G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC69305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes antisense oligonucleotides (AAX52869-X5571)

directed against at least 2 mRNAs selected from target genes, coding and

conn-coding regions of RNAs corresponding to target genes, gene initiation

codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'

codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'

codons, mixtures. The antisense oligonucleotides may be derived

from sequences AAX55272-74. These multiple target oligonucleotides

conditions or mixtures. The multiple target oligonucleotides

from sequences AAX55180-271) can be used for the antisense treatment of

diseases and conditions. Typical diseases and conditions are those

associated with impaired respiration and inflammation, including lung

diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis,

acute asthma, allergies, asthma, impeded respiration, respiratory

diseases syndrome, pain, cystic fibrosis, pulmonary hypertension,

pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary

disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g.

colon cancer, breast cancer, hung cancer, paincreatic cancer,

colon cancer, breast cancer, melanoma, hepatic metastasized

colon cancer, breast cancer, melanoma, hepatic metastasized

to the lungs, including breast and prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tumour necrosis factor alpha; TNF-alpha; antisense oligonucleotide; ASO; inhibition; expresssion; treatment; disease; disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides used in treatment of, e.g. pulmonary vasoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.4%; Score 14.8; DB 1; Length 21; Best Local Similarity 88.9%; Pred. No. 1.8e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumour necrosis factor alpha antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 6; Page 93; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1180 CGGCCCGGCTGACCCTG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX09096 standard; DNA; 21 BP
                                                                                                                                                                                            98WO-US019419
                                                                                                                                                                                                                                         97US-0059160P
98US-00093972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 cedeccedecreecee
                                                                                                                                                                                                                                                                                                                 (UYEC-) UNIV EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
prostate cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rattus rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUN-1999
                                                                                          WO9913886-A1
                                                                                                                                                                                         17-SEP-1998;
                                                                                                                                                                                                                                           17-SEP-1997;
                                                                                                                                                                                                                                                                  09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9901139-A1
                                                                                                                                            25-MAR-1999
                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX09096;
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Gaps

Pimstone SN;

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The invention relates to the human ABC1 cholesterol transporter protein

(B38082) and to nucleic acid sequences (C69120) which encode it. ABC1 is
a member of the ATP-binding cassette (ABC transporter) superfamily of
the ATP-binding cassette (ABC transporter) superfamily of
corrections, and plays a crucial role in cholesterol transport, particularly
convolved in cholesterol Efficiking in monocytes and fibroblasts, being
convolved in cholesterol efficik from the cell. The gene encoding ABC1 is
clocated on chromosome 9q31 and mutations in this gene encoding ABC1 is
convolved in cholesterol by and familial HDL deficiency (FHA). These diseases
convolved in cholesterol in the blood correlate with a high risk of cardiovascular
cholesterol in the blood correlate with a high risk of cardiovascular
disease, coronary restencesis, and peripheral vascular disease.
conversely, a high level of HDL has protective effects against
cardiovascular disease. The invention promptising human ABC1 nucleic
cardiovascular disease. The invention promptising human ABC1 nucleic
cardiovascular disease computing the administration of an expression
cerdiovascular disease comprising the administration of an expression
cardiovascular disease comprising whether a patient has an
cincreased risk for cardiovascular disease due to polymorphisms in the
corrector encoding ABC1 or an active fragment thereof. The invention also
correctored the for determining whether a patient has an
cincreased risk for cardiovascular disease due to polymorphisms in the
corrector encoding and methods of screening for such compounds
correctored the and an expression
corrector Human ABC1 proceins and nucleotides can be used to treat
cardiovascular disease, corrective disease, well and are processed risk for cardiovascular disease, and methods of screening for such co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention specifically excludes proteins with the exact amino acid esquences of Genbank Accession No: CAA10065.1 and X79926, and the nucleic acid with the exact sequence as Genbank Accession No: AJ012376.1. The present sequence represents a polymorphic site of the human ABC1 gene
                                                                                                                                                                                               New ABC1 polypeptide is useful for treating diseases associated with ABC1 biological activity, e.g. Alzheimer's disease, Huntington's disease and
                                                                                                                                                                                                                                                                                                                Example, Fig 11; 229pp; English.
UNIV BRITISH COLUMBIA
XENON BIORESEARCH INC.
                                                                                     Wilson AR,
                                                                                                                                       WPI; 2000-587528/55
                                                                                     Hayden MR,
                             (XENO-)
                                                                                                                                                                                                                                                          cancer
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ö 0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; Indels Sequence 21 BP; 2 A; 5 C; 10 G; 4 T; 0 U; 0 Other; 0; Mismatches 16; Conservative Query Match Best Local Similarity tches

514 1 ACACGCTGGGGGTGCTGG 18 497 ACACGCTGGACGTGCTGG g

AAA32322 standard; DNA; 21 AAA32322; RESULT 1752 AAA32322, MARKEX BX SX SX B

BP.

A2b adenosine receptor antisense oligonucleotide SEQ ID NO:10.

(first entry)

28-JUL-2000

Human, adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiallergor; antiallergic; antiallergic; cytostatic; analgesic; impaired airway;

lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain, cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.

WO200009525-A2.

24-FEB-2000

99WO-US017712. 03-AUG-1999;

98US-0095212P.

03-AUG-1998;

(UYEC-) UNIV EAST CAROLINA.

Nyce JW;

WPI; 2000-205971/18.

New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or cancers.

Example 5; Page 257; 1343pp; English.

The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%) which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antialial antial ergies, and/or antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation.

Consecul for the treatment of diseases and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. including lung diseases and diseases whose secondary condetions, pulmonary vasconstriction, allergies, asthma, confined respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (CODP), and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONS reduces side effects. The A-containing ONS break down with the concentration and inflammation. AAA32313 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present convention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 from the previously named sequences. SEQ ID NO:1 to 1680 (AAA32332 to AAA3392) are specifically claimed ONS from the present invention. N.B. Sequences given in the disclosure of the present invention in the disclosure of the present invention of the present intending the disclosure of the present invention. N.B. Sequences given in the disclosure of the present invention. N.B. with their corresponding SEQ ID NO: sequences given in the sequence

Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;

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Gaps

Gaps ö Score 14.8; DB 1; Length 21; Pred. No. 1.8e+03; 0; Mismatches 2; Indels 0.4%; 16; Conservative Local Similarity Query Match Best Loca Matches

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1180 CGGCCCGGCTGACCCTG 1197 20 CGGGCCCGGCTGGCCCGG ò g

AAA03716 standard; DNA; 21 AAA03716; AAA03716/C 1D AAA037. XX AC AAA037 XX DT 19-MAY XX

RESULT 1753

BP

(first entry) 19-MAY-2000 Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; bund usurfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pain; cystic fibrosis; allergic rhinitis; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;

Human adenosine A2b receptor antisense oligonucleotide #10.

(first entry)

14-MAR-2001

AAF18443;

AAF18443 standard; DNA; 21 BP.

RESULT 1754 AAF18443/c

1180 CGGCCCGGCTGACCCTG 1197

Local Similarity 88.9 nes 16; Conservative

Matches

20 cedececederiesceces

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Human adenosine A2b receptor antisense oligonucleotide SEQ ID NO:1000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA03715 represent specifically claimed phosphorothioate antisense oligonucleotides for use in the composition of the present invention. AAA02718, AAA02720, AAA02722 and AAA03716 to AAA03720 represent other phosphorothioate oligonucleotides used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel composition for treating or preventing e.g. cardiopulmonary and
                                     Human; adenosine Al receptor; antisense oligonucleotide; hypoxia; adenosine As receptor; adenosine As receptor; adenosine As receptor; phosphorothicate; cardioqualmonary failure; renal failure; ischaemia; endotoxin release; ARDS; acute respiratory distress syndrome; cytoprotective; anti-allergic; anti-inflammatory; anti-hypoxic; appraventricular tachycardia; allergic rhinitis; acute inflammation; chronic obstructive pulmonary disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8; DB 1; Length 21; Pred. No. 1.8e+03; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 52; 252pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%;
                                                                                                                                                                                                                                                                                                                                                        99WO-US012775
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                                                                                                                                                                                                                                                                                                                                                                                                                   98US-00093972
98US-0088657P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-116433/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce JW, Hill JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       renal injury.
                                                                                                                                                                                                             Ното варіелв.
                                                                                                                                                                                                                                                                        WO9963938-A2.
                                                                                                                                                                                                                                                                                                                                                        08-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                        09-JUN-1998;
                                                                                                                                                                                                                                                                                                                 16-DEC-1999.
                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Si
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Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers

and respiratory obstructions.

WPI; 2000-679539/66

Nyce JW;

24-MAR-2000; 2000WO-US008020. 06-APR-1999; 99US-0127958P. (UYEC-) UNIV EAST CAROLINA. (NYCE/) NYCE J W.

WO200062736-A2. Homo sapiens.

cancer; ss.

26-OCT-2000.

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The present invention describes low adenosine (A) content antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. C. (1) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities.

(I) can have respiratory bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities.

C. (I) can have respiratory disorders and malignancies, such as stimulating and expression and or activity of target polypeptides associated with corresponding peptide factors and transmitters, transcription factors, immunoglobuline and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, chemokine receptors, adenosine receptors, cytokines and chemokine receptors, adenosine receptors, brinding proteins, adenosine receptors, bradykinin receptors, cotokine and creaming respiratory obstruction and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders and receptors, binding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction) and/or lung inflammation, allergy (les) and/or surfactant hypoproduction which are associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, allergic fibrosis (RDS), pain, cystic fibrosis (CP), malmonary disease (CPDD), conditions between complements, infant prominers, and processes and medically pulmonary disease (CPDD), conditions and conditions elected from pulmonary vasoconstriction elected from pulmonary infantion elected from pulmonary vasoconservatory distress syndrome conditions elected from pulmonary vasoconservatory distress syndrome conditions and pulmonary infantions elected from pulmonary infantions elected from pulmonary infantions elected from pulmonary infantions elected f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 14; Page 306; 1592pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the present invention
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Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;

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Gaps ö

BP.

AAA99535 standard; DNA; 21

AAA99535/

19-JAN-2001

AAA99535;

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2213 AACAATGTGAGGGGTCCC 2230
                            19 AACAATGTCAGGGATCCC
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Gaps

; 0

2; Indels

0; Mismatches

16; Conservative

Matches

Best Local Similarity

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o;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The patent discloses Arabidopsis thaliana chloroplast targetted beta-amylase (ct beta-amylase) gene and a light or sugar inducible promoter sequence located on chromosome 4. Transgene comprising ct beta-amylase gene and the promoter sequence may be used to transform plants to alter the activity of starch biosynthesis or degradation pathway enzymes, target proteins or enzymes to plant plastids, alter seed set and impart herbicide or pest resistance. Manipulation of the amount of starch in plastids of leaves or storage organs, finds use in tobacco, potato crisp industry and others involving malting of grain, production of syrups and alcohol, paper making, pharmaceuticals, glue, oil, textiles and dairy products. The present sequence is the PCR primer P2, used for amplification of cDNA encoding A. thaliana chloroplast targetting transit peptide from ct beta-amylase gene located in cosmid AAG16599
                                                                                                                                                                                                                                                                                                                 Primer P2 to amplify A. thaliana chloroplast targetting transit peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel beta-amylase genes, useful for regulating starch in transgenic plants, has a chloroplast targeting sequence and new promoter sequence useful for the control of transgenic gene expression.
                                        Gaps
                                                                                                                                                                                                                                                                                                                                                  Chloroplast targetted beta-amylase; ct beta-amylase; promoter; chromosome 4; starch metabolism; plastid; herbicide resistance; pest resistance; syrup; alcohol; paper making; pharmaceutical; glue; textile; dairy product; chloroplast targetting transit peptide;
                                      ô
Score 14.8; DB 1; Length 21;
Pred. No. 1.8e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 4 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ADTE-) ADVANCED TECHNOLOGIES CAMBRIDGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 34; 74pp; English
                                                                        1180 CGGCCCGGCTGACCCTG 1197
                                                                                                         20 cedececederiedecede 3
                                                                                                                                                                                                      AAZ51371 standard; DNA; 21 BP
0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98GB-00017963.
99GB-00013014.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98GB-00017959
                                                                                                                                                                                                                                                                               (first entry)
                   Local Similarity 88.9
les 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kavanagh TA, Lao NT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-237643/20.
                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200011144-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .9-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-AUG-1998;
05-JUN-1999;
                                                                                                                                                                                                                                                                               06-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAR-2000
 Query Match ·
Best Local Sim
Matches 16;
                                                                                                                                                                                                                                          AAZ51371;
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                                                                                                                                                                    RESULT 1755
                                                                                                                                                                                       AAZ51371/c
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                                                                                                            Adenosine A2b receptor; phosphorothioate; validating target gene; pain satiety; anxiety; mood disorder; libido; cognition; sleep; cancer; respiratory condition; pulmonary condition; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determining the existence of a correlation between the pathology of a disease and a gene or mRNA encoding a target polypeptide suspected of being associated with the disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Pred. No. 1.8e+03;
0; Mismatches 2; Indels
                                                                               Adenosine A2b receptor mRNA antisense oligonucleotide V.
                                                                                                                                                                                                                                                                /*tag= a
/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 32; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1180 CGGGCCGGCTGACCCTG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA93732 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                             02-MAR-2000; 2000WO-US005643.
                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.98;
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                                               (first entry)
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Best Local Similarity 88.9
Matches 16; Conservative
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/*tag=
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                                                                                                                                                                                                                                                                                                              WO200051621-A1
                                                                                                                                                                                                                                Key
modified_base
                                                                                                                                                                                                Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nyce JW;
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ID AAA95
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vivlemore401-10.rng

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Ndael; Na+ anion driven exchanger gene; intracellular ion concentration, neurological disease; muscular disease; cardiac arrhythmia; hypertension; hypotension; angina; asthma; diabetes; renal insufficiency; urinary incontinence; acidosis; alkalosis; irritable colon; epilepsy; cerebrovascular ischaemia; autoimmune disease; gene therapy; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention is concerned with the isolation of a number of versions of the Drosophila Na+ anion driven exchanger NDAE1 coding sequence. They were isolated by sequencing expressed sequence tags and determining the function of the proteins. The NDAE1 protein is involved in the regulation of intracellular ion concentrations, and agonists and antagonists can be used in the treatment of a number of neurological and muscular diseases, as well a cardiac arrhythmias, hypertension, hypotension, angina, asthma, diabetes, renal insufficiency, urinary incontinence, acidosis, alkalosis, irritable colon, epilepsy, oreceptorvascular isochamia and autoimmune disease. The nddel gene can be used in the production of transgenic animals, such as knockout animals. Primers AAA93712-A93752 and AAA94200-A94222 were all used in the
                                                                       Drosophila Na+ driven ion exchanger gene ndael PCR primer SEQ ID NO: 40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New purified DNA useful in assays to detect the ability of agents which inhibit or enhance NDAE1 channel activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide array; genotyping; single base extension reaction; SBE; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sciortino CM, Shrode LD, Sussman CR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP flanking sequence #79 used in multiplexing PCR/SBE assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.8; DB 1; Length 21;
88.9%; Pred. No. 1.8e+03;
rative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 2 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isolation of sequences of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 26; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYCA-) UNIV CASE WESTERN RESERVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4
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                                                                                                                                                                                                                                                                                                                                                        17-FEB-2000; 2000WO-US004028
                                                                                                                                                                                                                                                                                                                                                                                             99US-0120620P
                                                                                                                                                                                                                                                                                                                                                                                                            16-FEB-2000; 2000US-00505647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2575 ACATCACAGGGTGCGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21 ACAGGACAGGGTGCGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAC73384 standard; DNA; 21
                                    12-JAN-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-565368/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                WO200049048-A1
                                                                                                                                                                                                                                           Drosophila sp.
                                                                                                                                                                                                                                                                                                                                                                                           18-FEB-1999;
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                                                                                                                                                                                                                                                                                                                     24-AUG-2000.
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 AAA93732;
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The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one such polymorphic locus used in the present invention. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to form the oligonucleotide array. Note: This sequence includes a SNP represented by the degenerate codon in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonuclectide array; genotyping; single base extension reaction; SBE; polymorphic locus; single nucleotide polymorphism; 88.
                                                                                                                                                                                                                                                                                              Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                     Hirschhorn JN, Huang X, Kaplan P, Lander ES, Lockhart DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.4%; Score 14.8; DB 1; Length 21; Best Local Similarity 80.0%; Pred. No. 1.8e+03; Matches 16; Conservative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 6 A; 7 C; 5 G; 2 T; 0 U; 1 Other;
                                                                                                                                                                       (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2121 CTCAGGGACGACTCCGTGT 2140
                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 56; 70pp; English.
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                                                                                          27-MAR-2000; 2000WO-US008069.
                                                                                                                         99US-0126473P.
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                                                                                                                                                                                                                                                                  WPI; 2000-656171/63.
                                                                                                                                                                                                                                     Sklar P;
                               WO200058516-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200058516-A2
 Unidentified.
                                                                                                                         26-MAR-1999;
23-JUN-1999;
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23-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-FEB-2001
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                                                            05-OCT-2000.
                                                                                                                                                                                                                                   Ryder T,
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WPI; 2001-226749/23
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                                                                                                                                                                     The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one such polymorphic locus used in the present invention. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to form the oligonucleotide array. Note: This sequence includes a SNP represented by the degenerate codon in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forenaics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                         Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                           Lockhart DJ;
                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mccarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                           Score 14.8; DB 1; Length 21; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                       3; Indels
                           Huang X, Kaplan P, Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gargill M, Ireland JS, Bolk S, Daley GQ,
                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 7 C; 5 G; 2 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human gene single nucleotide polymorphism #166.
                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                2121 CTCAGGGGACGACTCCGTGT 2140
                                                                                                                                               Example 7; Page 50; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                               1 CTCAGAGGACRACCCCGAGT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                                                                                                                                                                                                                                                                                                                                                             0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF95405 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 80.0'
Matches 16; Conservative
                          Hirschhorn JN,
(AFFY-) AFFYMETRIX INC
                                                                WPI; 2000-656171/63
                                      Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200118250-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
Variation
                                     Ryder T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mccarthy JJ;
                                     applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; \(\nu_{1}\); \(\nu_{2}\); \(
Nucleic acids comprising single nucleotide polymorphisms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 4 A; 7 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human gene single nucleotide polymorphism #522.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the human gene SNPS shown in the specification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                     Example; Page 59; 242pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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nes 16, Conservative
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                                                                                                                       atherosclerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF95761;
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                                                                                                                                                        The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thrombosmbolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a method for treating a patient diagnosed as having a lower than normal high density lipoprotein. cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, involving administering a compound that modulates Ltx. or RXR-mediated transcriptional activity or ABC1 expression or activity. The LXR gene product may be used in an assay to identify compounds useful for the treatment of a disease or condition selected a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High density lipoprotein-cholesterol; HDL-C; cardiovascular; ABC1; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.8; DB 1; Length 21;
88.9%; Pred. No. 1.8e+03;
iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 4 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wild type sequence for ABC1 polymorphic site #18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 4; 317pp; English.
                                                                                                 Example; Page 84; 242pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1309 GACGATGCCACTGACAAG 1326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAR-2000; 2000US-00526193.
23-JUN-2000; 2000US-0213958P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-MAY-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transcriptional activity.
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   analysis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   status of CpG islands within samples of genomic DNA. Sequences AAF22041 - AAF22094 represent PCR primers and probes specific for certain human gene fragments, used in examples illustrating the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting methylated CpG containing nucleic acids comprises treating DNA with sodium bisulfite to create methylation-dependent sequence difference, and detecting methylated DNA by fluorescence-based quantitative polymerase chain reaction.
                                                                                                                                                                                                                                                                                                                                                                                                                    Human; cytosine methylation; CpG island; methylation pattern detection;
                                                                                                                         Gaps
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lower than normal HDL cholesterol level, a higher than normal triglyceride level, and a cardiovascular disease
                                                                                   Score 14.8; DB 1; Length 21;
Pred. No. 1.8e+03;
0; Mismatches 2; Indels
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Pred. No. 1.8e+03;
0; Mismatches 2; Indels
                                                  Sequence 21 BP; 2 A; 5 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 3 A; 0 C; 11 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                     for human unmethylated ESR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Danenberg KD;
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                                                                                                                                                         497 ACACGCTGGACGTGCTGG 514
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                                                                                    0.4%;
ilarity 88.9%;
Conservative
                                                                                                                                                                                  1 ACACGCTGGGGGTGCTGG
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                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                    Reverse PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-025026/03
                                                                                                      Best_Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer; 88,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200070090-A1
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                                                                                                                                                                                                                                                                                                                                                   13-MAR-2001
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AAF22046
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vivlemore401-10.rng

agammaglobulinaemia; diaberes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabry's disease; familial hypercholesterolaemia; polycystic kidney disease; cancer; hereditary spherocytosis; polycystic kidney disease; cancer; hereditary spherocytosis; lengiectasia; familial colonic polyposis; bhlers-Danlos syndrome; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; inflammation; nervous system disorder; infection; rheumatoid arthritis; multiple sclerosis; diabetes; systemic lupus erythematosus; Graves disease; longevity; obesity; baldness; fertility; forensic; paternity testing; ss.

single nucleotide polymorphism; SNP; sickle cell anaemia;

Human single nucleotide polymorphism #37.

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cancer; head; neck;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a polynucleotide comprising an Skp 2 promoter sequence. The polynucleotide or a plasmid of vector comprising the polynucleotide is useful as a pharmaceutical, and in manufacturing a medicament for the treatment of cancer such as cancers of the breast head or neck, lung, stomach, colon, rectum, or uterine cervix. The promoter is useful in tumour and cancer therapy to kill or reduce the growth, division or viability of tumour or cancer cells. The present sequence is a mutant of mouse Skp2 promoter E2F second region binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polynucleotide comprising an Skp 2 promoter sequence, useful as a pharmaceutical and in manufacturing a medicament for the treatment of cancer such as cancers of the breast, head, neck, lung stomach, colon,
                                                                                                                  Mouse Skp2 promoter E2F second region binding site mutant #1.
                                                                                                                                                douse; Skp2 promoter; cytostatic; therapy; tumour; cancer.
lung; stomach; colon; rectum; uterine; cervix; mutant; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 5 A; 10 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NOVS ) NOVARTIS FORSCHUNGSSTIFTUNG ZWEIGNIEDERL.
                                                                                                                                                                                                                                       Location/Qualifiers
replace(7, G)
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                                                                                                                                                                                                                                                                      /*tag= a
replace(14, G)
                                                                                                                                                                                                                                                                                                /*tag= b
replace(17, G)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Krek W;
                           AAD27707 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rectum, or uterine cervix.
                                                                                      (first entry)
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                                                                                      18-APR-2002
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                                                                                                                                                                                             Mus sp.
Synthetic.
                                                         AAD27707;
                                                                                                                                                                                                                                                     mutation
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                                                                                                                                                 Mouse;
RESULT 1764
               4AD27707
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Nucleic acid comprising single nucleotide polymorphisms, useful in forensics, paternity testing and diagnosis of disease.

Claim 1; Page 39; 96pp; English.

Lander ES;

Cargill M, Ireland JS,

WPI; 2002-315108/35

(CARG/) CARGILL M. (IREL/) IRELAND J S. (LAND/) LANDER E S.

18-JAN-2001; 2001US-00765081. 19-JAN-2000; 2000US-0176861P.

US2002037508-A1. Homo sapiens.

28-MAR-2002.

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The invention relates to a nucleic acid comprising single nucleotide polymorphisms (SNPs) associated with diseases. The nucleic acids comprising the SNPs and probes and primers for detecting them may be used in assays for the diagnosis of diseases associated with SNPs (such as congrising the SNPs and probes and primers for detecting them may be used in assays for the diagnosis of diseases associated with SNPs (such as SICKIE cell anaemia, agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan SCC syndrome, muscular dystroppy, Wiskott-Aldrich syndrome, Fabry's disease, familial hypercholesterolaemia, polycystic kidney disease, hereditary spheroxytosis, Von Willebrand's disease, tuberous sclerosis, hereditary syndrome, osteogenesis imperfecta, and acute intermittent porphyria, syndrome, osteogenesis imperfecta, and acute intermittent porphyria, syndrome, osteogenesis imperfecta, and acute intermittent porphyria, symptoms of, or susceptibility to, multifactorial diseases of which a component is or may be genetic, such as autoimmune diseases of which a component is or may be genetic, such as autoimmune diseases of which a thritis, multiple sclerosis, diabetes (insulin-dependent and non-confidence), systemic lupus erythematosus and Graves disease, cancers including cancers of the bladder, brain, breast, colon, oesophagus, the stomach and unterus, longevity, and susceptibility or receptivity to particular drugs or therapeutic treatments), in forensics and in paternity testing. Absidness and in paternity testing, Absidness and in paternity testing, Absidness and in paternity testing. Absidness and in paternity testing and the bladder, breast treatments), in forensics and in paternity testing and the breast colon, desceptibility or and in paternity testing and the breast colon, desceptibility or the patricular drugs or therapeutic treatments), in forensics
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hes 16; Conservative
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Gaps

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Indels

0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03;

0; Mismatches

88.98;

Local Similarity 88.5 nes 16; Conservative

Best Loca Matches

Query Match

2717

2700 TCCCACCCTGCCCCTCAG 3 TCCCACCCGCCACCTCAG

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BP.

ABK65417 standard; DNA; 21

RESULT 1765

02-JUL-2002 (first entry)

ABK65417;

ABK65417/ ID ABK6 XX AC ABK6 XX DT 02-J

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Gaps

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Indels

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1; Mismatches

Matches 16; Conservative

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Human; single nucleotide polymorphism; SNP; sickle cell anaemia; agammaglobulinaemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabry's disease; familial hypercholesterolaemia; polycystic kidney disease; cancer; hereditary spherocytosis; Von Willebrand's disease; tuberous sclerosis; hereditary paemorrhagic telangisctasia; familial colonic polyposis; Ehers-Danlos syndrome; osteogenesis imperfects; autoimmune disease; acute intermittent porphyria; inflammation; nervous system disorder; infection; rheumatodia arthritis; multiple sclerosis; diabetes; systemic lupus erythematosus; Graves disease; longevity; obesity; baldness; fertility; forensic; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid comprising single nucleotide polymorphisms, useful in forensics, paternity testing and diagnosis of disease.
                                                                                                Human single nucleotide polymorphism #266.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lander ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 69; 96pp; English.
                                                                                                                                                                                                                                                                                                                                               18-JAN-2001; 2001US-00765081.
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                        ABK65646 standard; DNA; 21
                                                                        02-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                Cargill M, Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                               (CARG/) CARGILL M.
(IREL/) IRELAND J S.
(LAND/) LANDER E S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-315108/35
                                                                                                                                                                                                                                                                                                JS2002037508-A1
                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                ABK65646;
RESULT 1766
              ABK65646/c
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The invention relates to a nucleic acid comprising single nucleotide

polymorphisms (SNPs) associated with diseases. The nucleic acids

comprising the SNPs and probes and primers for detecting them may be used

in assays for the diagnosis of diseases associated with SNPs (such as

sickle cell anaemia, agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan

syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabry's disease,

familial hypercholesterolaemia, polycystic kidney disease, hereditary

syndrome, osteogenesis imperfects, and acute intermittent porphyria,

syndroms of, or susceptibility to, multifactorial diseases of which a

component is or may be genetic, such as autoimmune diseases,

component is or may be genetic, such as autoimmune diseases,

component is or susceptibility to, multifactorial diseases of which a

component is or susceptibility to, multifactorial diseases

component is or succeptibility to, and factorial diseases

component is or succeptibility to, and factorial diseases

component is or succeptibility to, and sucception by

pathogenic microorganisms, autoimmune diseases, cancers

inflammation, cancer, diseases of the nervous system, and infection by

pathogenic microorganisms, autoimmune diseases, cancers

including cancers of the bladder, brain, breast, colon, ossophagus,

conceptivity, leukaemia, liver, lung, oral cavity, ovary, pancreas, prostate,

skin, stomach and uterus, longevity, appearance (e.g., baldness,

creceptivity to particular drugs or therapeutic treatments), in forensics

and in paternity testing. ABK65381-ABK65841 represent human single

cup components of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 9 C; 7 G; 2 T; 0 U; 1 Other;
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed clones are cultured and the calculant cultures are amplified by using the above primer; (g) signals are detected from the amplified by using the above primer; (g) signals are detected from the amplified by using the above primer; (g) here are
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                                                                                                                                                                                                                                                                                      Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. AB142957 to AB145322 represent PCR primers for human chromosome 1036-35 DNA, and AB145323 to AB145634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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18.9%; Pred. No. 1.8e+03;
ve 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                  Human chromosome 1p36-35 PCR primer SEQ ID NO:1121.
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2082 GTACTCCCGGGTGGCCAGG 2101
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                     20 GTCCTGCCCRGGAGGCCAGG 1
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0.4%; Score 14.8; DB 1; Length 21; 80.0%; Pred. No. 1.8e+03;

Best Local Similarity

Query Match

ABX09457/c

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The invention relates to a multidrug resistance-associated protein 1 (MRP -1) polynucleotide. The polynucleotide is useful in an in vitro method for identifying as single nucleotide polymorphism and for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of MRP-1 or for identifying and obtaining an inhibitor of the activity of a molecular variant of MRP-1. The sequences are useful for diagnosing a disorder related to the presence of a molecular variant of MRP-1 or susceptibility to such a disorder, where the disorder is cancer (particularly renal cancer) or a disease related to multidrug resistance. This sequence represents a human MRP-1 polymorphic DNA region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel multidrug resistance-associated protein 1 polynucleotide useful for diagnosis and treatment of cancer and multidrug resistance related diseases, and for identifying single nucleotide polymorphisms.
                          Human; multidrug resistance-associated protein 1; MRP-1; ss; cancer; renal cancer; cytostatic; single nucleotide polymorphism.
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88.9%; Pred. No. 1.8e+03;
ative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                      26-JAN-2001; 2001EP-00101651.
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                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel method for determining the genetic risk of arteriosclerosis both for clinical diagnosis and for population studies. The method comprises: (i) selecting risk-associated reference nucleic acid sequences, including their functionally characterizing complements, to a carrier; (iii) hybridising the probes with a nucleic acid from (or synthesised from) a patient sample; and (iv) detecting and evaluating the hybridisation pattern. The method provides a quick, inexpensive and informative diagnosis, and makes possible a miltifactorial analysis for detecting e.g. synergism between different mutations or mutations that when present alone carry no risk but are risk associated in presence of other mutations. The results may be combined with known risk-assessment methods to provide a more reliable diagnosis, especially important with new therapeutic methods (e.g. gene therapy) that are directed against specific genes. All relevant mutations in a reference sequence can be screened for in a single test and the method is well suited to automation. Abs. 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determining genetic risk of arteriosclerosis, for clinical diagnosis, comprises hybridizing patient nucleic acid with an array of probes derived from risk-associated reference genes and their mutations.
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                                                                                                                                                                                                                                                 Arteriosclerosis; diagnosis; hybridisation; synergism; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.8; DB 1;
Pred. No. 1.8e+03;
0; Mismatches 2;
                                                                                                                                                                                          Arteriosclerosis-detecting probe from HNF1 #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human MRP-1 polymorphic DNA region #175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 126; 146pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2911 CATGGCCCTGGGCGGGC 2928
                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 ccreeccereceeeeec 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-MAR-2002; 2002WO-EP002780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-MAR-2001; 2001DE-01011925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%;
                    ABX09457 standard; DNA; 21
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                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.4
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seedorf U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-723374/78.
                                                                                                                                                                                                                                                                              mutation; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (OGHA-) OGHAM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                       WO200272882-A2
                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                   22-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                              19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cullen P,
                                                                           ABX09457;
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RESULT 171
ABS66910/1
ID ABS61
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AC ABS61
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DT 29-NC
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Gaps

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1) polymucleotide. The polymucleotide is useful in an in vitro method for identifying a single nucleotide polymorphism and for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of MRP-1 or for identifying and obtaining an inhibitor of the activity of a molecular variant of MRP-1. The sequences are useful for diagnosing a disorder related to the presence of a molecular variant of MRP-1 or susceptibility to such a disorder, where the disorder is cancer (particularly renal cancer) or a disease related to multidrug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antianginal; cardiant; vasotropic; antiatherosclerotic; vulnerary; cerebroprotective; antiulger; antiasthmatic; antiallergic; antimigraine; nootropic; tranquilliser; neuroleptic; antimanic; antidepressant; stroke; antiemetic; gene therapy; hypertension; urinary retention; osteoporosis; angina pectoris; myocardial infarction; restenosis; atherosclerosis; aneuryem; wound healing; ischaemia; ulcer; asthma; allergy; migraine; vomiting; benign prostatic hypertrophy; psychotic; neurological disorder; degenerative disease; PCR primer; ss.
                                                                                                                                       Novel multidrug resistance-associated protein 1 polynucleotide useful for diagnosis and treatment of cancer and multidrug resistance related diseases, and for identifying single nucleotide polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a multidrug resistance-associated protein 1 (MRP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             resistance. This sequence represents a human MRP-1 polymorphic DNA region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Movel human adenylate cyclase type IV protein, regulators of which are useful for treating and preventing atherosclerosis, hypertension, osteoporosis, ulcer, asthma, allergy, psychotic and neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; adenylate cyclase type IV; enzyme; hypotensive; osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human adenylate cyclase type IV PCR primer SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 8 A; 4 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1;
Pred. No. 1.8e+03;
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   Mornhinwed E;
                                                                                                                                                                                                                                                                                          Example 2; Page 78; 198pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2791 TACATTTCTATAAATAGA 2808
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-OCT-2000; 2000US-0241306P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.98;
Hoffmeyer S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABN89299 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floeckner J, Liu N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-507943/54.
                                                                       WPI; 2002-657475/70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200233100-A2
   Brinkmann U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-APR-2002
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The present invention describes human adenylate cyclase type IV (I). (I) has hypotensive, osteopathic, antianginal, cardiant, vasocropic, antiatherosclerotic, vulnerary, cerebroprotective, antiulcer, antiallerotic, concluding the treatment of an adenylate cyclase type IV dysfunction related disease, in particular hypertension, urinary retention, osteoporosis, angina pectoris, myocardial infarction, ceretensists, atherosclerosis, a disease characterised by excessive smooth muscle cells or reduced smooth muscle cell proliferation, aneurysms, wound healing stroke, ischaemia, ulcer, asthma, allergy, benign corder, including anxiety, schizophrenia, manic depression, depression, delirium, dementia, severe mental retardation and degenerative diseases. The present sequence represents a PCR primer for amplifying the adenylate cyclase type IV gene from a human endothelial cell cDNA library in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a /*tag= https://withchistory.etrand.coverhangs/hote= "the 3'end of this strand by the sequence 5'-UU-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New method for the synthesis of target-specific short double stranded RNAs by hybridizing the sense oligonucleotide product with the complementary antisense oligoribonucleotide product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid synthesis; small interfering RNA; siRNA; RNA polymerase; promoter; T7; transcription; EGFP; green fluorescent protein; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a new method for the synthesis of target-
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                                                                                                                                                                                                                                                                                                                                                                                                                         21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1; Length
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                   Seguence 21 BP; 2 A; 11 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
Example 2; Page 53; 95pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Fig 3A; 35pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EGFP target-specific siRNA 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-OCT-2002; 2002WO-EP012165.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-NOV-2001; 2001US-0337975P.
                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     868 GAGGCTGACGAGGCGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 GGGCTGAAGAGGCGGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 88.5.
Local Similarity 88.5.
Local Similarity
Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (JANC ) JANSSEN PHARM NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    De Backer MD, Harris AN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1772
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Gaps

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Length 21; 2; Indels

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combining a target-specific sense or antisense collocurication and a chain extending enzyme in a reaction mixture so,that the template extended sense or antisense oligonucleotide product is formed; and (b) hybridizing the sense oligonucleotide product with the complementary antisense oligoribonucleotide product. The target-specific ds RNAs are useful in RNA interference in vertebrate and invertebrate systems as small interfering RNAs (siRNAs). The method provides truncated RNA polymerase promoter sequences, where the substitutions do no affect the in vitro transcription yields, but increase the possibility that at least one target-specific sequence of a defined sequence length exists in the RNA of the target protein. Sequence ACCY8430-431 represent BGFP target-specific ds siRNAs used in a FACS analysis of EGFP-transfected HeLa cells
specific short double stranded (ds) RNAs. The method involves:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 3 A; 5 C; 8 G; 0 T; 5 U; 0 Other;
85000000000000558
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Ouery Match 0.4%; Score 14.8; DB 1; Length 21; Best Local Similarity 72.2%; Pred. No. 1.86+03; 2; Indels 3; Mismatches 1530 CGAGGAGCAGCTCACCTT 1547 CGAGGAGCUGUUCACCUU 21 Conservative 13; Matches

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Gaps

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ADA15942 standard; DNA; 21 BP ADA15942; Synthetic storage protein oligonucleotide SM91.

(first entry)

06-NOV-2003

ss; lysC; transgenic; lysine accumulation; dihydrodipicolinic acid synthase; DHDPS; lysine inhibition; lysine ketoglutarate reductase; LKR; chloroplast transit sequence; CTS; aspartokinase III; AKIII; synthetic seed storage protein; SSP.

Synthetic.

US6459019-B1.

01-OCT-2002

24-MAR-1997;

97US-00823771

92US-00855414. 94US-00178212. 95US-00474633. 19-MAR-1992; 36-JAN-1994; 07-JUN-1995;

(DUPO) DU PONT DE NEMOURS & CO

Falco SC, Keeler SJ, Rice JA;

WPI; 2003-028272/02.

Transformed plants that accumulate lysine at higher levels in its seeds than untransformed plants, has gene fragments encoding lysine-insensitive dihydrodipicolinic acid synthase and lysine ketoglutarate reductase.

Example 21; Col 79; 109pp; English.

The invention relates to a plant comprising two foreign nucleotide sequences which cause seeds obtained from the plant to accumulate lysine at a level of at least 10% higher than seeds of a plant that do not comprise the nucleotide, where the nucleotide comprises a fragment encoding a dihydrodipicolinic acid synthase (DHDPS) that is insensitive to lysine inhibition, and a fragment encoding a plant lysine ketoglutarate reductase (LKR) or its subfragment. The nucleotide fragment is operably linked to a plant chloroplast transit sequence (CTS) and the plant lysine ketoglutarate reductase subfragment is used in antisense ADA15942/C
ID ADA15942/C
XX AC ADA1594
XX ADA1594
XX BS; lyi
XW BS; lyi
XW BS; lyi
XX BYDHE
XX BYDHE
XX BYDHE
XX BYDHE
XX BYD OI-OCT
XX BYDHE
XX BYDH
XX BYDHE
XX BYDH

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above mentioned plant and seeds obtained from the above mentioned plant. The seeds obtained from the above mentioned plant. The seeds obtained from the above mentioned plant (e.g., rapeseed, soybean or corn) comprising the foreign nucleic acid sequences accumulate lysine at a higher level, preferably at a level of at least 10% higher than seeds of a plant that do not comprise the foreign nucleic acid asquences. Chimaeric gene comprising DHDPS from C. glutamicum and aspartchinase III (from the lysC gene) of E. coli (mutated to be lysine-insensitive) are also used to generate the above transgenic plants. Also disclosed are synthetic seed storage proteins (SSP) used as an internal source of lysine, built up from synthetic peptide monomers based around an Earl site sequence (for generating mutlimeric proteins). The present sequence is a strand of an oligonucleotide encoding an SSP monomer.
inhibition or cosuppression. Also included are progeny plants from the
   $$9$$$$$$$$$$$$$$$$
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Sequence 21 BP; 2 A; 9 C; 0 G; 10 T; 0 U; 0 Other;

Gapa . 0 0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; Live 0; Mismatches 2; Indels Local Similarity 88.9 nes 16; Conservative Query Match Best Loca Matches

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ACH03698 standard; DNA; 21 BP. ACH03698; ACH03698/c

RESULT 1774

Ear I-based lysine-rich heptad repeat oligonucleotide SM91. (first entry) 25-SEP-2003

Aspartokinase; AKIII; dihydrodipicolinic acid synthase; DHDPS; seed lysine content; seed threonine content; seed storage protein; SSP; chloroplast transit sequence; lysine-rich protein; lysine ketoglutarate reductase; LKR; transgenic; ss.

Synthetic

US2003056242-A1.

20-MAR-2003.

17-DEC-2001; 2001US-00023066.

93WO-US002480. 94US-00178212. 95US-00474633. 97US-00823771. 92US-00855414. 19-MAR-1992; 18-MAR-1993; 06-JAN-1994 07-JUN-1995 24-MAR-1997

s C (FALC/) FALCO

Falco SC;

WPI; 2003-521869/49.

New nucleic acid fragment encoding aspartokinase and dihydrodipicolinic acid synthase, useful for increasing threonine or lysine content of seeds acid synt.

Example 21; Page 43; 116pp; English.

The invention relates to an isolated nucleic acid fragment comprising a first nucleic acid subfragment encoding aspartokinase (AK) that is substantially insensitive to inhibition by lysine, and a second nucleic acid subfragment encoding dihydrodipicolinic acid synthase (DHDPS) that is substantially insensitive to inhibition by lysine. Also included are an isolated nucleic acid fragment comprising a nucleic acid subfragment encoding lysine ketoglutarate reductase (LKR), a chimaeric gene (where

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the mucleic acid fragment is operably linked to a plant chloroplast transit sequence and to a seed-specific regulatory sequence, a plant comprising the mucleic acid/chimaeric gene in its genome, a seed obtained from the plant, increasing threonine or lysine content of the seeds of plant having the ability to produce levels of free threonine or lysine at least two times greater than the free threonine levels of untransformed least two times greater than the free threonine levels of untransformed least two plants, a transformed (soybean) plant comprising seeds that accumulate lysine at a level at least two percent to four-fold higher than the seeds of an untransformed rapesed comprising seeds that accumulate lysine to a level between ten percent and one hundred percent of opportising in its genome the nucleic acid fragment having the monocot-comprising in its genome the nucleic acid fragment having the monocot-comprising in its genome the nucleic acid fragment having one hundred that the percent higher than the seeds of the untransformed plant. Also thirty percent higher than the seeds of the untransformed plant. Also cisclosed are synthetic lysine-rich seed storage proteins (SSP), built up from monomer lysine-rich seed storage proteins (SSP), built up from monomer lysine-rich seeds to a pool of lysine in a transformed plant. The nucleic acid fragments, genes and methods are useful for increasing threonine or lysine content of the seeds of the need to supplement mixed grain feeds with lysine or threonine produced via microbial fermentation. The present sequence is one strand of a DNA contain a produced via a monomer unit in a monomer produced are surfaced produced are surfaced are produced are surfaced are produced and menced and menced are surfaced are contained broader are sequence is one strand of a DNA contain a post and methods are sequenced and menced and produced are surfaced at the plant is and methods are not in a produced and microbial stransformed between the plant in a produced and microbial stransfor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 2 A; 9 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     synthetic seed storage protein
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Gaps ö 0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; 2; Indels 0; Mismatches Local Similarity 88.9 es 16; Conservative Query Match tches

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ADC53895 standard; DNA; 21 18-DEC-2003 ADC53895 RESULT 1775 ADC53895/

BP

Pediococcus dextrinicus detection PCR primer #3. (first entry)

Pediococcus bacteria detection; spacer region; 16S rRNA; 23S rRNA; beer brewery; fermented-food; 88; PCR; primer.

Pediococcus dextrinicus.

JP2003038181-A.

12-FEB-2003

01-JUN-2001; 2001JP-00165989

01-JUN-2001; 2001JP-00165989.

(ASAK) ASAHI BREWERIES LTD.

WPI; 2003-590691/56

Oligonucleotide for detecting bacteria, comprises sequence of spacer region between genes encoding 16S rRNA and 23S rRNA of Pediococcus dextrinicus, Pediococcus parvulus, Pediococcus acidilactici, Pediococcus pentosaceus

Claim 9; SEQ ID NO 11; 18pp; Japanese

The invention comprises oligonucleotides for detecting bacteria, the

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method for detecting or quantitating a nucleic acid analyte in a sample comprising contacting the sample with an analyte-binding oligonucleotide, a labelled oligonucleotide, a capture polymer and a linear stem oligonucleotide under conditions where a complex is formed comprising the analyte, analyte-binding oligonucleotide, labelled oligonucleotide, capture polymer and linear stem oligonucleotide; and detecting or quantitating the complex. The method is useful in detecting or quantitating a nucleic acid analyte in a sample. The present sequence represents a probe for human FC mRNA, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
oligonucleotides contain a gene sequence of a spacer region between genes encoding 16S rRNA and 23S rRNA of Pediococus dextrinicus, P. parvulus, P. acidilactici or P. pentosaceus. The oligonucleotides of the invention are useful as primers for detecting P. dextrinicus, P. parvulus, P. acidilactici and P. pentosaceus, in beer brewery and in fermented-food preparation process. The present DNA sequence represents a detection PCR primer of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting or quantitating a nucleic acid analyte in a sample by contacting the sample with an analyte-binding, labeled and linear stem oligonucleotide and a capture polymer to form a complex and detecting o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detection; quantitation; nucleic acid analyte;
analyte-binding oligonucleotide; capture polymer; human; haemoglobin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                        Score 14.8; DB 1; Length 21;
Pred. No. 1.8e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a probe fin an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 21 BP; 0 A; 9 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                              Sequence 21 BP; 9 A; 3 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6; SEQ ID NO 135; 94pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Fc mRNA probe SEQ ID NO:135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wong WLT,
                                                                                                                                                                                                                                                        3562 ACTGCTACCTTTCAAAGC 3579
                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-MAR-2003; 2003WO-US009726.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-2002; 2002US-0368669P.
                                                                                                                                                                                Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                         18 ACTGCTTCTTTTCAAAGC
                                                                                                                                                                                                                                                                                                                                                                                 ADD01591 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               quantitating the complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tsai SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-804090/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003083440-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stephan JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                    ADD01591;
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                                                                                                                                                                                                                                                                                                                                               RESULT 1776
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1180 CGGCCCGGCTGACCCTG 1197

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation coden, coding regain, 5 or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway definication and a second active agent comprising an entilnflammatory steroid and ubjquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine or receptor, producing bronchodilation, increasing levels of definitions or receptor, producing bronchodilation, increasing levels of butterior is the composition or condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lung surfactant in a subject's tissue, or treating bronchoconstriction, ung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.4%; Score 14.8; DB 1; Length 21; Best Local Similarity 88.9%; Pred. No. 1.8e+03; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 9379; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human phosphorothioate antisense fragment,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
1522 AAGCCGCCCGAGGAGCAG 1539
                                                                                                                                                                                                                                                                           BP.
                                                                          18 AAGCCGCGGAAGGAGCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                           ABZ94137 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                    17-OCT-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229219/22.
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ubiquinone
                                                                                                                                                                                                                                                                                                                                                          ABZ94137;
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XX AB29
XX A
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The invention relates to a method for generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells. The method comprises exposing the precursor cells or exocrine cells to: a nucleic acid molecule encoding neurogenin 3 (ngn3) or NeuroD/Deta2; or an activator of ngn3 or NeuroD/Deta2 gene expression, under conditions effective to generate the insulin-generating cells from the precursor or exocrine cells. The invention is useful in generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells is useful for generating glucose sensitive insulin secreting beta cells useful for generating of or in situ development of insulinsecreting cuitable for transplantation, and for in situ development of insulinsecreting premature differentiation of precursor stem cells into insulin-secreting premature differentiation of precursor stem cells into insulin-secreting cell differentiation. The present sequence is human RT-PCR primer for isolation of mouse ngn3 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                         stem cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells, for generating glucose sensitive insulin secreting beta cells for transplantation, comprises using neurogenin3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
                                                                                                                                                      Human; insulin-secreting cell; neurogenin 3; ngn3; precursor st
pancreatic exocrine cell; transplantation; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1; Length 21; Pred. No. 1.8e+03; 0; Mismatches 2; Indel8
                                                                                                                      Human RT-PCR forward primer for mouse ngn3 DNA isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 7 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5B; Page 37; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                Gradwohl G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               489
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ABD32529/c
1D ABD32529 standard; DNA; 21 BP.
XX
               AAD47326 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                         26-FEB-2002; 2002WO-DK000130.
                                                                                                                                                                                                                                                                                                                                                          26-FEB-2001; 2001US-0271474P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.48;
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                                                                                      24-FEB-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 88.9
                                                                                                                                                                                                                                                                                                                                                                                          (NOVO ) NOVO NORDISK AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Heimberg H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-018804/01.
                                                                                                                                                                                                                                                 WO200274946-A2
                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NeuroD/beta2.
                                                                                                                                                                                                                                                                                     26-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Serup P,
                                                AAD47326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Matches
AAD47326,
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Gaps

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers; (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition
                                                                                                                                          Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respirator, distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the invention has antiallergic, and intinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted unucleic acids associated with lung airway or lung dysfunction, and
                                                                                                 A1/A3 antisense receptor-associated oligonucleotide SEQ ID 9379.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 1 A; 9 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; SEQ ID NO 9379; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A, K.
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC
                                                29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bronchodilating agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                                                                                                                                                      WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nyce JW, |
Miller S,
  ABD32529;
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Aguilar D;

Pabalan J,

Katz.E,

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periodontal ligament fibroblast and its use for detection of periodontal ligament fibroblast and its use for detection of periodontal ligament fibroblast, mass-isolation of PDLs17 protein, and production of an antibody specific to the PDLs17. The specification also discloses a polypeptide encoded by an mRNA is selected from the group consisting of PDLs5, PDLs21, PDLs22, PDLs31 which are selectively expressed in periodontal ligament fibroblast and a biological probe selected from cDNAs of PDLs 5, 17, 22, 25 and 31. An antigenic peptide for PDLs17 is selected from the peptides disclosed in the specification. An antibody capable of binding the PDLs17 polypeptide is prepared by administrating the PDLs17 peptide into an animal. This sequence corresponds to a PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polypeptide pdls17 specific to periodontal ligament fibroblast and the use thereof.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel PDLs17 polypeptide which is specific to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                            Gaps
                                                                                                                                                                                                                                                                            88; periodontal ligament fibroblast; PDLs5; PDLs17; PDLs22; PDLs25; PDLs31; primer.
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Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                      Periodontal ligament fibroblast gene PDLs cDNA primer #2.
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                          Indels
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 9 A; 4 C; 3 G; 5 T; 0 U; 0 Other;
                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer to amplify a PDLs cDNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 13; 31pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Park JC;
                                                        1180 CGGCCCGGCTGACCCTG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3568 ACCTITCAAAGCTIGGAG 3585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 ACCTTTCAAACATGGAG 18
                                                                                   m
                                                                                                                                                                BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                        04-APR-2001; 2001KR-00017905.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-APR-2001; 2001KR-00017905
          88.9%;
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                                                                                                                                                              ADP19689 standard; DNA; 21
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                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 88.9
                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-338954/32.
       Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PARK/) PARK J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Han GY, Kim HJ,
                                                                                                                                                                                                                                                                                                                                                            KR2002078533-A
                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                        12-AUG-2004
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                                                                                                                                                                                          ADP19689;
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                                                                                                                             RESULT 1780
ADP19689
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ADK98157/c
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0.4%; Score 14.8; DB 1; Length 21;

Query Match

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The invention relates to a novel predictor set comprising a plurality of polynucleotides and/or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein tyrosine kinase activity or members of the protein tyrosine kinase activity or members of the protein tyrosine pathway. The molecules of the invention demonstrate cytostatic, antiangiogenic, vasotropic and vulnerary activities and may be useful in the field of pharmacogenomics, in particular for determining drug sensitivity and in treating breast cancer, hypervascular diseases, angiogenesis and scars in wound healing. The current sequence is that of a human protein tyrosine kinase biomarker-related RT-PCR primer of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytostatic; Antiinflammatory; Immunosuppressive; Antibacterial; Virucide; cancer; inflammatory; immune; ds; human secreted protein.
                                      polypeptides whose expression pattern predicts cell response to a compound that modulates protein tyrosine kinase activity, useful in
                  New predictor sets with a plurality of polynucleotides and/or
                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted protein encoding sequence SEQ ID #1192.
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 9 A; 7 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                               Disclosure; SEQ ID NO 311; 649pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2316 TCTGTGTGTGTGTGTG 2333
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2002US-0406585P.
2002US-0406588P.
2002US-0406608P.
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2002US-0406642P.
2002US-0406646P.
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2002US-0406655P.
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2002US-0410946P.
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2002US-0410948P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 16; Conservative
                                                                        treating breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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29-AUG-2002;
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ADP29194/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 predictor set; protein tyrosine kinase biomarker; cytostatic; antiangiogenic; vasotropic; vulnerary; pharmacogenomic; drug sensitivity; breast cancer; hypervascular disease; angiogenesis; wound healing scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human protein tyrosine kinase biomarker-related RT-PCR primer SEQ ID 311.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                    Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single nucleotide polymorphism in human gene.
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88.9%; Pred. No. 1.8e+03;
ve 0; Mismatches 2; Indels
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                                                    human; single nucleotide polymorphism; SNP; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 7 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; SEQ ID NO 7186; 2627pp; Japanese.
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                                                                                                                                                                                                                                                                                           (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2338 TGTGTGTGTGTGCACA 2355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; ss; RT-PCR; PCR; primer.
                Primer of the invention #3877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADL61387 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                     08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                              08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-AUG-2003; 2003WO-US026491
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nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                  WPI; 2004-093977/10.
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                                                                                                                                 JP2003259875-A.
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                                                                                                                                                                        16-SEP-2003.
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                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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RESULT 1782

ADL61387,

Best Loc Matches

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Gaps

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Gaps

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Indels

0.4%; Score 14.8; DB 1; ilarity 88.9%; Pred. No. 1.8e+03; Conservative 0; Mismatches 2;

Query Match Best Local Similarity Matches 16; Conserv

Length 21;

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New nucleic acid molecule for diagnosing, preventing or treating diseases such as proliferative (e.g. cancer), inflammatory, immune, metabolic, genetic, bacterial and viral diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to an isolated nucleic acid molecule ancoding a polypeptide which is believed to be cytostatic, antiinflammatory, immunosuppressive, antibacterial and virucidal. The composition and methods are useful for diagnosing, preventing and treating diseases such as proliferative (e.g. cancer), inflammatory, immune, metabolic, genetic, bacterial and viral diseases. The present sequence represents a human secreted brotein encoding sequence. The present sequence is available on WIPOWEB and is not in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Williams LT, Chu K, Lee E, Hestir K, Beaurang PA, Behrens D
Halenbeck RF, Huang MM, Kothakota S, Haishan L, Linnemann T;
Pierce K, Wang Y, Wong JGP, Wu G, Zhang H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (FIVE-) FIVE PRIME THERAPEUTICS INC.
                        17-SEP-2002; 20020S-0410961P.
17-SEP-2002; 20020S-0410962P.
17-SEP-2002; 20020S-0411019P.
17-SEP-2002; 20020S-0411022P.
17-SEP-2002; 2002US-0411023P.
17-SEP-2002; 2002US-0411024P.
17-SEP-2002; 2002US-0411035P.
17-SEP-2002; 2002US-0411035P.
17-SEP-2002; 2002US-0411035P.
17-SEP-2002; 2002US-0411035P.
17-SEP-2002; 2002US-0411041P.
                                                                                                                                                                                                           17-SEP-2002; 2002US-0411046P.
17-SEP-2002; 2002US-0411046P.
17-SEP-2002; 2002US-0411048P.
17-SEP-2002; 2002US-0411052P.
17-SEP-2002; 2002US-0411052P.
17-SEP-2002; 2002US-041101P.
17-SEP-2002; 2002US-041101P.
17-SEP-2002; 2002US-041101P.
18-APR-2003; 2003US-0463700P.
18-APR-2003; 2003US-0463700P.
18-APR-2003; 2003US-0463700P.
18-APR-2003; 2003US-046370P.
18-APR-2003; 2003US-046370P.
18-APR-2003; 2003US-046370P.
                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAY-2003; 2003US-0467239P.

02-MAY-2003; 2003US-0467230P.

19-MAY-2003; 2003US-0471306P.

19-MAY-2003; 2003US-0471336P.

22-MAY-2003; 2003US-0472420P.

22-MAY-2003; 2003US-0472430P.

09-JUN-2003; 2003US-047669P.

09-JUN-2003; 2003US-0476641P.

08-JUL-2003; 2003US-0485213P.

08-JUL-2003; 2003US-0485213P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUL-2003; 2003US-0485325P.
14-JUL-2003; 2003US-0486446P.
14-JUL-2003; 2003US-0486480P.
15-JUL-2003; 2003US-0486891P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-AUG-2003; 2003US-0493573P
08-AUG-2003; 2003US-0493577P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nuclectide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                breast cancer; cytostatic; gene therapy; human; chromogranin B; CHGB; secretogranin 1; SCG1; chromosome 20pter-p12; ss; PCR; primer; SNP; single nucleotide polymorphism.
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                                                                                                                                                                                                                     Extend primer 37 used to genotype human chromogranin B polymorphism.
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1221 CTTCGGCCAGGTGGTCAT 1238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2306 AGAGCTTTGGTCTGTG 2323
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 AGAGCTTTCATCTGTGTG
                                                                                                                  ADP09242 standard; DNA; 21
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ADP48125
ID ADP48125
XX
AC ADP48125
                                                                               RESULT 1784
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Behrens D;

Sequence 21 BP; 8 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

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20-APR-1999;
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25-0CT-2000
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                                                                                                                                        Kurane R,
Kovama O,
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                                                                                                                                                     Koyama
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                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a novel isolated polynucleotide comprising a nucleic acid sequence, the human MRCK1 gene located at position 11q13, and the novel human protein kinase MRCK1 encoded by it. The sequence of the invention has sequence homology to rat myotonic dystrophy kinase-related Cdc42 binding kinase (MRCK). The invention may be useful for disagnosing, prognosing, and treating kinase-related diseases, preferably diseases associated with aberrant expression of MRCK1. The present sequence is that of a DNA sequence which is part of the human MRCK1 gene which may be a target for a short inhibitory (siRNA) sequence and which is related to the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                           New human protein kinase MRCKI with 65% sequence homology to rat myotonic dystrophy kinase-related Cdc42-binding kinase (MRCK), useful in diagnostics and as a drug target.
                                                               protein kinase; MRCK1; myotonic dystrophy kinase; Cdc42 binding kinase; MRCK; kinase-related disease; short inhibitory RNA; siRNA; ds; human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 1.8e+03; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                       Human MRCK1 siRNA target DNA sequence SeqID160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic deoxyribooligonucleotide poly i.
                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 160; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2063
                                                                                                                                                                                           07-NOV-2003; 2003WO-US035609
                                                                                                                                                                                                                     27-NOV-2002; 2002US-0429381P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 CGAGGAGTACCTGTACCT
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           (first entry)
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Best Local Similarity 86...
-hea 16; Conservative
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                                                                                                                                     WO2004050831-A2
                                                                                                                                                                                                                                                               (LIUW/) LIU W. (WULL/) WU L.
                                                                                                                                                                                                                                                 (AMHP ) WYETH.
                                                                                                                                                                                                                                                                                                        Wu L;
           09-SEP-2004
                                                                                                           Homo sapiens
                                                                                                                                                                 17-JUN-2004.
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Liu W,

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The invention relates to the determination of the concentration of a nucleic acid target, using a fluorescently labeled probe which produces creduced fluorescence emission when hybridised to the target nucleic acid. The method comprises measuring the reduction in emission caused by the real-time polymerase chain reaction, e.g. for nucleic acids by a real-time polymerase chain reaction, e.g. for quantifying microbial cells in co-oultures or symbictic systems, for quantifying microbial cells in co-oultures or symbictic systems, for detecting gene mutations or polymorphisms, and for analysing melting curves of target nucleic acids to determine a Th value. Methods of the invention allow target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and no accurately. Particularly there is no need to remove unbound probe, and no conventional PCR conditions can be used). The specificity of PCR is kept high (amplification of primer dimers is delayed), and the limit of conventional person of primer dimers is allowed, and amplification of primer dimers is allowed. Can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation coefficient than conventional graphs so more accurate quantitation is possible. The current sequence represents a synthetic characteristy of a target nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                              Yokomaku T;
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                                                                                                                                                                                                                                                                                                                                                                              Yamada K,
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                                                                                                                                                                                                                                                                                                                                                                    Kanagawa T, Kamagata Y, Kurata S,
Furusho K;
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                                                                                                                                                                               NIBI-) JAPAN BIOINDUSTRY ASSOC
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20-APR-2000; 2000EP-00108643.
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Measurement of nucleic acids, using a nucleic acid probe and analysis of
the obtained data.
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                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; fluorochrome; nucleic acid probe; fluorescence.
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                                                                                          (BIOI-) BIOINDUSTRY KYOKAI SH.
(KANK-) KANKYO ENG KK.
(KEIZ-) KEIZAI SANGYOSHO SANGYO GIJUTSU SOGO KEN
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(KEIZ-) KEIZAI SANGYOSHO SANGYO GIJUTSU SOGO KEN.
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24-AUG-1999; 99JP-0023666.
30-AUG-1999; 99JP-00242693.
01-FEB-2000; 2000JP-00028896.
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                       30-AUG-1999; 99JP-00242693.
01-FEB-2000; 2000JP-00028896.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to the determination of the concentration of a nucleic acid target, using a fluorescently labeled probe which produces reduced fluorescence emission when hybridised to the target nucleic acid. The method comprises measuring the reduction in emission caused by the method comprises measuring the reduction in emission caused by the method comprises measuring the reduction in emission caused by the real-time polymerase chain reaction, e.g. for charlifying microbial cells in co-cultures or symbiotic systems, for detecting gene mutations or polymorphisms, and for analysing melting curves of target nucleic acids to determine a Tm value. Methods of the invention allow target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and no materials are introduced that inhibit amplification by Taq polymerase (so conventional PCR conditions can be used). The specificity of PCR is kept of migh is reduced. Complex probes are not needed, and amplification can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation coefficient than conventional graphs a synthetic quantitation is possible. The current sequence represents a synthetic quantitation is decorrelation.
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                                                                                                                                                                                                                                                                                                                      Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
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                                                                                                                                                                                            Kanagawa T, Kamagata Y, Kurata S, Yamada K,
Purusho K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; fluorochrome; nucleic acid probe; fluorescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
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                                                                                          (NIBI-) JAPAN BIOINDUSTRY ASSOC.
(AGEN ) AGENCY OF IND SCI & TECHNOLOGY.
(KANK-) KANKYO ENG CO LTD.
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20-APR-2000; 2000EP-00108643.
                                                 99JP-00111601
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                                                                                                                                                                                                 Kurane R,
Koyama O,
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Query Match

Best Loca Matches

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RESULT 1788 ABA97620

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Gaps

RESULT 1791

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fluorescently-labeled nucleic acid probes for assaying nucleic acids and their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yamada K;
                                                                                                                                                                                                                                                                                                                                                    Probe; polymorphism detection; mutation detection; disease diagnosis; microbial identification; ss.
                                                                              Gaps
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                                             Score 14.8; DB 1; Length 30; Pred. No. 2.4e+03;
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                                                                              7; Indels
               Sequence 30 BP; 4 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kamagata Y, Torimura M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
                                                                           0; Mismatches
                                                                                                           3259 AGATATTTTATTTGCTTTGTCCTTTT 3284
                                                                                                                                                                                                                                                                                                                        Probe poly c for assaying nucleic acids
                                                                                                                                           Example 12; Page 60; 152pp; Japanese.
                                                                                                                                                                                                                          ABL95887 standard; DNA; 30 BP
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03-AUG-2000; 2000JP-00236115
26-SEP-2000; 2000JP-00292483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-JUN-2001; 2001WO-IB001147
                                           ch 0.4%;
1 Similarity 73.1%;
19; Conservative
                                                                                                                                                                                                                                                                                        (first entry)
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                                                            Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified.
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                                                                                                                                                                                                                                                          ABL95887;
                                             Query Match
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X 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-etranded oligonucleocide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and
                                                                                                                                                                                                                                                                                                                                                                                                                                          Yamada K;
                                                                                                                                               Probe, polymorphism detection; mutation detection; disease diagnosis;
microbial identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fluorescently-labeled nucleic acid probes for assaying nucleic acids their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Triple helix forming nucleotides 4791-4725 of the n-myc gene
                                                                                                                                                                                                                                                                                                                                                                                                                                          Kurata S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 0.4%; Score 14.8; DB 1; Length 30; 11 Similarity 73.1%; Pred. No. 2.4e+03; 19; Conservative 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 30 BP; 4 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Torimura M,
                                                                                                                                                                                                                                                                                                                                                                                         (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY (KANK-) KANKYO ENG CO LTD.
                                                                                                               Probe poly I for assaying nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3259 AGATATTTTATTTGCTTTGTCCTTTT 3284
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 12; Page 60; 152pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Kanagawa T, Kamagata Y,
                  BP.
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                                                                                                                                                                                                                                                                                                                        27-JUN-2000; 2000JP-00193133.
03-AUG-2000; 2000JP-00236115.
26-SEP-2000; 2000JP-00292483.
                                                                                                                                                                                                                                                                                             27-JUN-2001; 2001WO-IB001147.
                ABL95893 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX14632 standard; DNA; 35
                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         medicine for e.g. analyti
microbial identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-195876/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                             WO200208414-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the invention
                                                                                                                                                                                             Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-MAR-1999
                                                                               19-JUN-2002
                                                                                                                                                                                                                                                             31-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Kurane R, |
Yokomaku T;
                                                 ABL95893;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2
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Matches
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ID AAX1
XX AAX1
AC AAX1
XX DT 24-M
XX XX
ABL95893
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Gaps

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3259 AGATATTTTATTTGCCTTTT 3284

19; Conservative

Matches

à 셤

Best Local Similarity

5 Aratrititrititicrititrititi 30

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The present sequence represents a potential triple-helix forming region. It can be used to demonstrate the assay of the invention. The assay comprises adding a sample containing double-stranded DNA test sequences. e.g. containing the present sequence, to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is sequence. Triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                        Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
Triple-helix forming region; Triplex formation; DNA detection;
                identification; bacteria; oncogene; virus; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Col 13-14; 168pp; English.
                                                                                                                                                                                                                                                         (PROF-) PROFILE DIAGNOSTIC SCI INC.
                                                                                                                                                                          93US-00173489.
                                                                                                                                                                                                                 92US-00968436.
                                                                                                                                                                                                                                                                                                Hepburn AG, Wang C;
                                                                                                                                                                                                                                                                                                                                     WPI; 1999-130384/11.
                                                         Homo sapiens
                                                                                                                                                                                                                   29-OCT-1992;
                                                                                              US5861244-A
                                                                                                                                     19-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                         bacteria.
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Gaps .. 0 DB 1; Length 35; Query Match 0.4%; Score 14.8; DB 1; Length 3 Best Local Similarity 64.7%; Pred. No. 2.6e+03; Matches 22; Conservative 0; Mismatches 12; Indels Sequence 35 BP; 1 A; 4 C; 0 G; 30 T; 0 U; 0 Other;

3300 TTCTATAGGATTTTTCTTTAGGAGATTTATTTT 3333 1 rrrcrarrcrirrrrrrrrrrrrrrrrrrrrrr ઠે d

AAS95728 standard; DNA; 45 BP AAS95728; RESULT 1793 AAS95728

14-FEB-2002 (first entry)

Allele discrimination Pl primer #12.

Rolling circle amplification; single nucleotide polymorphism; anaemia; exonuclease deficient DNA polymerase; amplification target circle; RCA; Parkinson's disease; polycystic kidney disease; Tay-Sachs disease; ss; Huntington disease; sickle cell anaemia; haemophilia; cystic fibrosis; diabetes; obesity; cancer; head; neck; skin; brain; oesophagus; stomach; nung; breast; colon; ovary; testis; prostate; leukaemia; lymphoma; melanoma; PCR primer; sequencing primer; probe.

Homo sapiens

WO200177390-A2

18-OCT-2001

The invention relates to detecting single nucleotide polymorphisms by contacting an allele-specific oligonucleotide primer (PI) with a target contacting an allele-specific oligonucleotide primer (PI) with a target contacting an allele-specific oligonucleotide primer (PI) with a target sequence is complementary to PI at one end but the terminal nucleotide and the circumal nucleotide and the complementary. The complex is then contacted with an exchuclease complementary. The complex is then contacted with an excurclease deficient DNA polymerase enzyme under conditions that promote extension of PI with the target DNA as the template, thereby forming an extended segment of PI. Oligonucleotide probes whyridising to one or more target polymucleotides distinguish between matched and mismatched 3' ends, hence the absence of sequence amplification indicates the presence of a single amplification target circle can be used in rolling circle amplification can be used in rolling circle amplification and is useful for diagnosing a disease caused by, induced by or related to a mutation in at least one gene, such as parkinson's disease, polycystic kidney disease, Tayancha disease, Huntington disease, sickle cell anaemia, haemophilia, cystic fibrosis, diabetes, obesity, cancers of the head, neck, skin, brain, oseophagus, stomach, lung, breast, colon, ovary, testis or prostate, leukaemia, lymphoma and malanoma. Sequences AASSS711-AASSS745 represent primers, targets and fluorescence decorators used in the detection of RCA products Detecting single nucleotide polymorphism involves amplifying target sequences using small primer probe that matches or mismatches to target sequence and extending primer probe which is then detected. Claim 15; Page 41; 67pp; English (MOLE-) MOLECULAR STAGING INC 05-APR-2000; 2000US-0194843P. 05-APR-2001; 2001WO-US011151 WPI; 2002-049157/06. Abarzua P;

ö Query Match 0.4%; Score 14.8; DB 1; Length 45; Best Local Similarity 73.1%; Pred. No. 2.8e+03; Matches 19; Conservative 0; Mismatches 7; Indels Sequence 45 BP; 2 A; 5 C; 0 G; 38 T; 0 U; 0 Other;

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Gaps

3262 TATTTTATTTGCTTTGTCCTTTTTCA 3287 12 rrrrrrrrrrrrrrrrrrrrrrrrrrrr

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AAS95724 standard; DNA; 45 AAS95724; RESULT 1794 AAS95724

BP

Allele discrimination Pl primer #8.

14-FEB-2002 (first entry)

Rolling circle amplification; single nucleotide polymorphism; anaemia; exonuclease deficient DNA polymerase; amplification target circle; RCA; Parkinson's disease; polycystic kidney disease; Tay-Sackh disease; se; Huntington disease; sickle cell anaemia; haemophilia; cystic fibrosis; diabetes; obesity; cancer; head; neck; skin; brain; oesophagus; stomach; lung; breast; colon; ovary; testis; prostate; leukaemia; lymphoma; melanoma; PCR primer; sequencing primer; probe.

Homo sapiens.

WO200177390-A2

18-OCT-2001.

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The invention relates to detecting single nucleotide polymorphisms by contacting an allele-specific oligonucleotide primer (PI) with a target contacting an allele-specific oligonucleotide primer (PI) with a target contacting an allele-specific oligonucleotide and the polymorleotide to form a hybridisation complex, where the target sequence is complementary to PI at one end but the terminal nucleotide and the third nucleotide from the terminal at the other end of PI may not be complementary. The complex is then contacted with an excondlease deficient DNA polymerase enzyme under conditions that promote extension of PI with the target DNA as the template, thereby forming an extended complex is 10 oligonucleotide probes hybridising to one or more target polymucleotide given matched and mismatched 3' ends, hence the absence of sequence amplification indicates the presence of a single conclectide mismatch. Primer sequences complementary to a sequence on an amplification target circle can be used in rolling circle amplification amplification in at least one gene, such as Parkinson's disease, polycystic kidney disease, Tay-Sachs disease, Huntington disease, sickle cell anaemia, haemophilia, cystic fibrosis, diabetes, obesity, cancers of the head, neck, skin, brain, osephagus, stomach, lung, breast, colon, ovary, testis or prostate, leukaemia, lymphoma and melanoma. Sequences AAS95711-AAS95745 represent primers, targets and contacts in the detection of RCA products
                                                                                                                                                                                                                                                                                          Detecting single nucleotide polymorphism involves amplifying target sequences using small primer probe that matches or mismatches to target sequence and extending primer probe which is then detected.
                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; Page 41; 67pp; English
                                                                            05-APR-2000; 2000US-0194843P.
                                                                                                                                (MOLE-) MOLECULAR STAGING INC
                           05-APR-2001; 2001WO-US011151.
                                                                                                                                                                                                                                        WPI; 2002-049157/06.
                                                                                                                                                                                    Abarzua P;
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Sequence 45 BP; 1 A; 6 C; 0 G; 38 T; 0 U; 0 Other;

DB 1; Length 45; 7; Indels 2.8e+03; Score 14.8; DB Pred. No. 2.8e+ 0; Mismatches 3262 TATTTTATTTGCTTTGTCCTTTTTCA 3287 Query Match
Best Local Similarity 73.1%;
Matches 19; Conservative

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Gaps

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12 Trititititititititititititica 37 AAS95724 standard; DNA; 45 a

Allele discrimination P1 primer #8.

14-FEB-2002 (first entry)

AAS95724;

Rolling circle amplification; single nucleotide polymorphism; anaemia; exonuclease deficient DNA polymerase; amplification target circle; RCA; Parkinson's disease; polycystic kidney disease; Tay-Sachs disease; ss; Huntington disease; sickle cell anaemia; haemophilia; cystic fibrosis; diabetes; obesity; cancer; head; neck; skin; brain; oesophagus; stomach; lung; breast; colon; ovary; testis; prostate; leukaemia; lymphoma; melanoma; PCR primer; sequencing primer; probe.

Homo sapiens,

WO200177390-A2

18-OCT-2001

05-APR-2001; 2001WO-US011151.

05-APR-2000; 2000US-0194843P.

(MOLE-) MOLECULAR STAGING INC

WPI; 2002-049157/06.

Detecting single nucleotide polymorphism involves amplifying target sequences using small primer probe that matches or mismatches to target sequence and extending primer probe which is then detected.

Claim 15; Page 41; 67pp; English.

The invention relates to detecting single nucleotide polymorphisms by contacting an allele-specific oligonucleotide primer (PI) with a target contacting an allele-specific oligonucleotide primer (PI) with a target contacting an allele-specific oligonucleotide primer a polymucleotide and the complementary to PI at one end but the terminal nucleotide and the tird nucleotide from the terminal at the other end of PI may not be complementary. The complementary is then contacted with an expanclease of PI with the target DNA as the template, thereby forming an extended segment of PI oligonucleotide probes hybridising to one or more target oplymucleotides distinguish between matched and mismatched 3' ends, hence the absence of sequence amplification indicates the presence of a single nucleotide mismatch. Primer sequences complementary to a sequence on an amplification target circle can be used in rolling circle amplification amplification target circle can be used in rolling circle amplification of the method is useful for diagnosing a disease, such as Parkinson's disease, polycystic kidney disease, Tay-Sachs disease, Huntington

CH or related to a mutation in at least one gene, such as Parkinson's disease, sickle cell anaemia, haemophilia, cystic fibrosis, diabetes, obesity, cancers of the head, neck, skin, brail, long, breast, colon, ovary, testis or prostate, leukaemia, lymphoma and melanoma. Sequences ASSE711-AASSE715 represent primers, targets and fluorescence decorators used in the detection of RCA products

Sequence 45 BP; 1 A; 6 C; 0 G; 38 T; 0 U; 0 Other;

Gaps .; 0 Score 14.8; DB 1; Length 45; Pred. No. 2.8e+03; 0; Mismatches 17; Indels 0.48; Query Match 0.4 Best Local Similarity 59.5 Matches 25; Conservative

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1353 GGAGATGATGAAGATGATCGGGAAACACAAAAACATCATCAA 1394 ð 셤

ADD69515 standard; DNA; 15 BP RESULT 1796 ADD69515

ADD69515;

15-JAN-2004 (first entry)

ISSR-related PCR primer 2.

inter-simple sequence repeat; ISSR; SSR; PCR; primer; genotyping; plant; animal; Basmati fice; ss.

Unidentified

WO2003085133-A2.

16-OCT-2003.

09-JAN-2003; 2003WO-IB000041.

08-APR-2002; 2002IN-CH000260.

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Gaps

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New

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retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer. The agents can be used to treat prostate, breast, cervical and ovarian cancers and to make hormonal drugs for retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer. The drug can retard the change of hormone-dependent cancers into hormone-independent cancers effectively. The present sequence represents a PCR primer which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a hormonal drug-containing agent (I) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hormone dependent cancer; hormone independent cancer; hormonal drug; prostate cancer; breast cancer; cervical cancer; ovarian cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hormonal drug-containing agents for retarding conversion of horm dependent cancers, useful e.g. treating prostate and breast cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.6; DB 1; Length 17;
Pred. No. 1.5e+03;
                                                        Length 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                 Indels
                 Sequence 15 BP; 0 A; 0 C; 7 G; 7 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 2 A; 3 C; 6 G; 1 T; 0 U; 5 Other;
                                                      Score 14.6; DB 1;
Pred. No. 1.3e+03;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                  Human epidermal growth factor sense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 19; 31pp; Japanese.
                                                                                                 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1618 CACAGGGACCTGGCTGC 1634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
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                                                                                                                                                                                                                                                                                         ВР
                                                                                                                                          2319 GTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98JP-00286793
                                                        ch 0.4%; ll Similarity 93.3%; l4; Conservative
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                                                                                                                                                                                                                                                                                         AAA29596 standard; DNA; 17
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CAYMGGGAYYTGGCHGC
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CHEM IND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matsutani E, Naito K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-303644/26
                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hormone dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TAKE ) TAKEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200020034-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                           10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-APR-2000
                                                                                                                                                                                                                                                                                                                                  AAA29596;
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AAA29598
ID AAA29598
XX
AC AAA29598
                                                                                                                                                                                                                                                    RESULT 1798
                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                    셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New set of inter-simple sequence repeats (ISSR)-PCR primers for genotyping eukaryotes, useful for genotyping diverse genomes of plant and animal systems.
                                                                                                                         set of inter-simple sequence repeats (ISSR)-PCR primers for otyping eukaryotes, useful for genotyping diverse genomes of plant and
                                                                                                                                                                                                                                                             (ISSR)-PCR primers for genotyping eukaryotes. The primers of the invention may be useful for genotyping diverse genomes of plant and animal systems, in particular for distinguishing Basmati rice varieties from non-Basmati rice varieties and traditional Basmati rice varieties from evolved Basmati rice varieties. The current sequence is that of the ISSR-related PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel set of inter-simple sequence repeats (ISSR)-PCR primers for genotyping eukaryotes. The primers of the invention may be useful for genotyping diverse genomes of plant and animal systems, in particular for distinguishing Basmati rice varieties from non-Basmati rice varieties and traditional Basmati rice varieties from evolved Basmati rice varieties. The current sequence is that of the ISSR-related PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence repeat; ISSR; SSR; PCR; primer; genotyping; plant;
                                                                                                                                                                                                                                                    invention relates to a novel set of inter-simple sequence repeats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.6; DB 1; Length 15; 93.3%; Pred. No. 1.3e+03; ive 1; Mismatches 0; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 BP; 0 A; 0 C; 7 G; 7 T; 0 U; 1 Other;
DNA FINGERPRINTING & DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNAF-) CENT DNA FINGERPRINTING & DIAGNOSTICS
                                                                                                                                                                                                           Disclosure; Page 19; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 19; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2317 CTGTGTGTGTGTG 2331
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ISSR-related PCR primer 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     animal; Basmati rice; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-804317/75
                                                                                API; 2003-804317/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003085133-A2
                                                                                                                                                                    animal systems.
CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inter-simple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
                                      Nagaraju JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nagaraju JG
                                                                                                                                            genotyping
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(DNAP-)
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RESULT 1797

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ADD69514

IID ADD6

XX ADD6

YOU ADD7

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Matches

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Gaps

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This invention relates to a novel therapeutic agent for the treatment of hormone dependent cancer that comprises a hormone group chemical agent and a differentiation inducing agent. Specifically, the hormone group chemical agent refers to an agonist derived from the lutainising brownereleasing hormone (LH-RH), whereas the differentiation inducing agent is preferably a fat soluble vitamin, or can be a nuclear receptor ligand, histone acetylation regulation drug or a DNA methylation regulation drug or a DNA methylation regulation drug or therapeutic agent for hormone dependent prostate, ovarian, uterine or breast cancer, and can also delay change in hormone independent cancer. Accordingly, the compositions of the invention are described as degenerate PCR primer used to amplify the insulin receptor, in an exemplification of the invention.
                                                                                                                                                                       Novel therapeutic agent of hormone dependent cancer, comprising hormone group, chemical agent and differentiation inducing agent, useful for treating hormone dependent cancer in mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hormone dependent cancer; hormone; differentiation inducing agent; luteinising hormone-releasing hormone; LH-RH; fat soluble vitamin; prostate; ovarian; uterine; breast cancer; hormone independent cancer; anti-tumour; PCR; primer; ss; EGF receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Degenerate sense PCR primer used to amplify the EGF receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.6; DB 1; Length 17; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 2 A; 4 C; 4 G; 1 T; 0 U; 6 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6; Mismatches
                                                                                                                                                                                                                                                                Example 1; Page 14; 15pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1618 CACAGGGACCTGGCTGC 1634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 CAYMGRGACYTKGCWGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADH23250 standard; DNA; 17 BP
31-MAY-2002; 2002JP-00160837.
                                         31-MAY-2002; 2002JP-00160837.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-MAY-2002; 2002JP-00160837.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAY-2002; 2002JP-00160837
                                                                                  (TAKE ) TAKEDA CHEM IND LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-113108/12.
                                                                                                                                WPI; 2004-113108/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JP2004002240-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH23250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1801
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a hormonal drug-containing agent (1) for retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer. The agents can be used to treat prostate, breast, cervical and ovarian cancers and to make hormonal drugs for retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer sancer. The drug can retard the change of hormone-dependent cancers into hormone-independent cancers into a prement sequence represents a PCR primer which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hormone dependent cancer; hormone; differentiation inducing agent; luteinising hormone-releasing hormone; LH-RH; fat soluble vitamin; prostate; ovarian; uterine; breast cancer; hormone independent cancer; anti-tumour; PCR; primer; ss; insulin receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                       Hormone dependent cancer; hormone independent cancer; hormonal drug; prostate cancer; breast cancer; cervical cancer; ovarian cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hormonal drug-containing agents for retarding conversion of hormone-dependent cancers, useful e.g. for treating prostate and breast cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Degenerate sense PCR primer used to amplify the insulin receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.6; DB 1; Length 17; 54.7%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 17 BP; 2 A; 4 C; 4 G; 1 T; 0 U; 6 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 19; 31pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1618 CACAGGGACCTGGCTGC 1634
                                                             Human insulin sense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ||::||:|||:|||
CAYMGRGACYTKGCWGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH23252 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                            98JP-00286793.
                                                                                                                                                                                                                                                                                                                                 99WO-JP005533.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          64.78;
                                                                                                                                                                                                                                                                                                                                                                                                                      TAKE ) TAKEDA CHEM IND LTD
                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matsutani E, Naito K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-303644/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                            WO200020034-A1
                                                                                                                                                  PCR primer; 88
                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                 07-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                            08-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                    10-AUG-2000
                                                                                                                                                                                                                                                                                     13-APR-2000
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Query Match

Matches

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Gaps

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Novel therapeutic agent of hormone dependent cancer, comprising hormone group, chemical agent and differentiation inducing agent, useful for treating hormone dependent cancer in mammal.

JP2004002240-A

Synthetic

ADH23252;

RESULT 1800 ADH23252

08-JAN-2004.

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AAQ30386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                            hormone dependent cancer that comprises a hormone group chemical agent and a differentiation inducing agent. Specifically, the hormone group chemical agent cancer that comprises a hormone group chemical agent refers to an agonist derived from the luteinising hormone releasing hormone (IH-RH), whereas the differentiation inducing agent is preferably a fat soluble vitamin, or can be a nuclear receptor ligand, histone acetylation regulation drug or a DNA methylation regulation drug. The present invention describes using the IH-RH agonist as a preventative or therapeutic agent for hormone dependent proseate, ovarian, uterine or breast cancer, and can also delay change in hormone independent cancer. Accordingly, the compositions of the invention are described as exhibiting high anti-tumour activity. This oligonucleotide sequence is a degenerate PCR primer used to amplify the epidermal growth factor (EGF) receptor, in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acids, useful for regulating T-cell mediated immune responses, e.g., suppressing T-lymphocytes in subjects with autoimmune disorders, or enhancement in those with viral infections, comprises novel
                                                                     This invention relates to a novel therapeutic agent for the treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated nucleic acid which includes a promoter sequence being transcriptionally functional in a T-lymphocyte undergoing activation and transcriptionally less functional in the T-lymphocyte prior to the activation. The nucleic acid is useful for regulating T-cell mediated immune responses in mammals. Nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T-lymphocyte modulator; autoimmune disorder; graft rejection; PCR; graft-versus-host disease; viral infection; lymphocyte kinase; LCK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fugu rubripes lymphocyte kinase (LCK) amplifying PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.6; DB 1; Length 17; 70.6%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 2 A; 3 C; 6 G; 1 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MOLE-) INST MOLECULAR & CELL BIOLOGY. (EHRL/) EHRLICH G.
                        Example 1; Page 13; 15pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 21; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tan YH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1618 CACAGGACCTGGCTGC 1634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ||::||||::||||:||
1 CAYMGGGAYYTGGCHGC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              70.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-AUG-2001; 2001WO-IL000765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-SEP-2000; 2000US-0229326P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD34155 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brenner S, Venkatesh B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           I-cell active promoters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-329781/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Takifugu rubripes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200218619-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-AUG-2003
16-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD34155;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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AAD34155/
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molecules of the invention may be used to suppress or eliminate T-
lymphocytes undergoing activation to suppress T-lymphocyte mediated
immune response in individuals suffering from immune disorders, e.g.
autoimmune disorders such as graft rejection or graft-versus-host
disease. They may also be used to enhance T-lymphocyte mediated immune
response in individual suffering from, e.g. viral infection. The present
sequence is Fugu rubripes lymphocyte kinase (LCK) amplifying PCR primer
used in the exemplification of the invention. (Updated on 29-AUG-2003 to
standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumour necrosis factor, herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligomer TNF217 for forming triplex with HUMTNFAA target duplex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= b
/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= c
/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= f
/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                     tch 0.4%; Score 14.6; DB 1; Length 20; al Similarity 70.0%; Pred. No. 1.8e+03; 14; Conservative 2; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     methyl-8-oxo-2'
                                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 5 C; 5 G; 4 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                       1750 AAGTGGATGGCGCCTGAGGC 1769
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/mod_base= OTHER
/note= "OTHER= N6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'mod_base= OTHER
'note= "OTHER= N6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                 20 AARTGGACNGCNCCNGARGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= g
/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ30386 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ30386;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1803
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95US-00437465. 96US-00643411. 96US-00721327.

26-SEP-1996;

06-MAY-1996;

09-MAY-1995;

97US-00850961. 94US-00240030.

05-MAY-1997; 09-MAY-1994;

US6013517-A.

Synthetic.

11-JAN-2000.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human turnour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV, HER; HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448s. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gag; pol; retroviral vector construct; gag/pol expression cassette; antivaral; immunomodulatory; cytotoxin; prodrug activator; replacement gene; antisense sequence; ribozyme; tumour prevention; viral infection; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis, herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide #1 used in gag-pol expression cassette construction.
note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
0
                                                                                                                /mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine'
                                                                methyl-8-oxo-2' deoxyadenine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.6; DB 1; Length 21; 81.0%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                               Froehler B, Krawczyk S, Matteucci MD, Milligan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 11 A; 0 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3458 AAGTITATATATATCTATATA 3478
                                                   /mod_base= OTHER
/note= "OTHER= N6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 AAAATTATATATATATTA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 12; Page 70; 77pp; English.
                              /*tag= j
mod_base= OTHER
                                                                                                                                                                                                                                                                            910S-00643382.
910S-00683420.
910S-0068544.
910S-00686546.
910S-00686547.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ90067 standard; DNA; 22 BP.
                                                                                                                                                                                                                                91WO-US008811
                                                                                                                                                                                                                                                                90US-00617907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17; Conservative
                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                (GILE-) GILEAD SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
               modified_base
                                                                               modified base
                                                                                                                                                                                                                                                                                                                            17-APR-1991;
17-APR-1991;
27-SEP-1991;
                                                                                                                                                                 WO9209705-A1
                                                                                                                                                                                                                                25-NOV-1991;
                                                                                                                                                                                                                                                                                 18-JAN-1991;
08-APR-1991;
                                                                                                                                                                                                11-JUN-1992
                                                                                                                                                                                                                                                                23-NOV-1990
                                                                                                                                                                                                                                                                                                                17-APR-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ90067;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1804
AAZ90067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents an oligonucleotide used in the construction of gag-pol expression cassettes. The invention relates to a retroviral vector construct which consists of a 5'-long terminal repeat (5'-LTR); a tRNA binding site; an origin of second strand DNA synthesis; a 3'-LTR and gag/pol sequences modified to contain two or more stop codons. The invention also relates to a gag/pol expression cassette, and an env expression cassette. The retroviral construct has anticancer, antiviral and immunomodulatory activity. The retroviral constructs are used to produce recombinant retroviral particles for use in gene transfer, particularly gene therapy, e.g. to deliver heterologous sequences that encode cytotoxins, prodrug activators, replacement genes, antisense sequences or ribozymes, immune accessory molecules and viral immunogens, particularly for treatment or prevention of tumours, viral infections and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOMIN's Moloney murine leukaemia virus; mouse; retroviral backbone; LTR; gag/pol expression cassette; gag; pol; env; integrase; gene therapy; ss; tumour; cancer; viral infection; immune response; autoimmune response; graft rejection; cytostatic; antiviral; immunostimulant; PCR; primer; immunosuppressive; murine leukaemia virus 4070A amphotropic envelope; bovine growth hormone polyadenylation sequence; long terminal repeat.
                                                                                                                                                                                                                                                                                                                                                                New retroviral construct, used to produce retroviral particles for gene therapy, containing a gag/pol sequence that includes at least two stop codons, incapable of producing replicable virus by recombination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                              Respess JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 14.6; DB 1; Length 22; 81.0%; Pred. No. 1.9e+03; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                              Bodner M, Driver DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gag/pol expression cassette construction primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 9 A; 3 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2828 ATACATATATATATACAT 2848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 ATATATATATCGATACCAT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Col 24; 63pp; English.
                                                                                                                                                                                                                                                                                                Sauter S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK33880 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17; Conservative
                                                                                                                                                                                                                                                                                              Depolo NJ, Chada S,
                                                                                                                                                                                                                                                                                                                               WPI; 2000-159877/14.
                                                                                                                                                                                                                                                          (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus sp.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK33880;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1805
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US6333195-B1

99US-00263959 94US-00309335 95US-00531241

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WPI; 2004-059052/06
                                                                                                                                                  Hood LE, Rowen L;
                                                                                                           (HOOD/) HOOD L E.
                                                                                                                         (ROWE/) ROWEN L.
 US2002150891-A1.
                                                 05-MAR-1999;
                                                                         19-SEP-1994;
                                                                                     19-SEP-1995;
                        17-OCT-2002
 ö
                                                                                                                                                                                                                                                                                                The invention relates to a gag/pol expression cassette comprising a promoter, a gag/pol gene (I) and a polyadenylation sequence in which the 5' end of (I) has been modified to contain codons that are degenerate for gag, or the 3' end of (I) has been deleted without affecting the biological activity of the encoded integrase. The expression cassette and similar cassettee that express env protein, are used to produce recombinant retroviral particles by homologous recombination. These particles are gene transfer vectors, particularly for gene therapy of tumours or viral infections, also to induce an immune response, to treat or prevent diseases, or to suppress graft rejection or immune/autoimmune responses. This sequence represents an oligonuclectide primer used in construction of gag/pol expression cassettes of the invention
                                                                                                                                                                                                            New gag/pol expression cassette, for preparing retroviral particles for gene therapy, comprises a promoter, a gag/pol gene, and a polyadenylation sequence, and cannot form a replication competent virus by homologous recombination.
                                                                                                                                                              Driver DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.6; DB 1; Length 22;
                                                                                                                                                              Bodner M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 9 A; 3 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                              Chada S, Sauter S,
                                                                                                                                                                                                                                                                           Example 3; Col 24; 63pp; English.
                                                                      95US-00437465.
96US-00643411.
96US-00721327.
97US-00850961.
                                    07-JAN-2000; 2000US-00479776
                                                                                                                                                              Respess JG, Depolo NJ,
                                                                                                                                     (CHIR ) CHIRON CORP.
                                                                                                                                                                                      WPI; 2002-163181/21.
                                                                       09-MAY-1995;
06-MAY-1996;
26-SEP-1996;
                                                            09-MAY-1994;
                                                                                                           05-MAY-1997;
            25-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Gaps ö 4; Indels Local Similarity 81.0%; Pred. No. 1.9e+03; tes 17; Conservative 0; Mismatches 4 Matches

ADH70330 standard; DNA; 27 BP 25-MAR-2004 (first entry) ADH70330; RESULT 1806 ADH70330/c

Human Vbeta gene repeat sequence #120.

human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; arrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infections disease; viral infection; HIV; fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer; breast cancer; ds

Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.

CD40L poly-A tract sequence SEQ ID NO:9.

Homo sapiens

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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each 'Deta gene,' UberaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases including autoimmune disease, degenerative nervous system diseases, and neoplastic diseases. Autoimmune diseases include Addison's disease, arophic gastrits. Degenerative nervous system diseases include multiple arophic gastritis. Degenerative nervous system diseases include multiple clerophic astritis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple in hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those caused by viruses such as HIV, fungal infections such as those caused by chistogomes, filteria and bartail infections such as those caused by schistosomes, filteria and bacterial infections such as those caused by schistosomes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 14.6; DB 1; Length 27;
11.0%; Pred. No. 2.3e+03;
ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 27 BP; 23 A; 0 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                           Disclosure; SEQ ID NO 524; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3264 TTTTATTTGCTTTGTCCTTTT 3284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 81.0
Matches 17; Conservative
                                                                                                Vbeta gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            셤
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The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (1) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding antirheusatic, immunosuppressive and antirhilammatory activities, and can be used in gene therapy. (I) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                              New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 29 BP; 22 A; 4 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                         (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.
                                                                                                                                                                                                                                                                                                                              Example 1, Fig 3, 90pp; English.
                                        13-SEP-2000; 2000WO-US024966.
                                                                                99US-0153625P
                                                                                                                                                                                                       WPI; 2001-244776/25.
                                                                                                                                                               Crow MK, Li Y;
                                                                                13-SEP-1999;
  22-MAR-2001
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0.4%; Score 14.6; DB 1; Length 29; 69.0%; Pred. No. 2.4e+03; 9; Indels 3304 ATAGGATTTTTCTTTAGGAGATTTATTTT 3332 0; Mismatches 20% Conservative Local Similarity Matches 8 셤

Query Match

AAF74924 standard; DNA; 29 BP RESULT 1808
AAF74924/C
XX
AC AAF74924
AC A

AAF74924;

23-MAY-2001 (first entry)

CD40L poly-A tract sequence SEQ ID NO:21

Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.

Homo sapiens

WO200119844-A1.

22-MAR-2001

13-SEP-2000; 2000WO-US024966

(NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.

99US-0153625P

13-SEP-1999;

Crow MK, Li Y;

WPI; 2001-244776/25

New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as

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The present invention describes an isolated, purified nucleic acid, which residues 331-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (I) has antiarthritic, and can be used in gene therapy. (I) is useful in the study, diagnosis and can be used in gene therapy. (I) is useful in the study, diagnosis and which elevated expression of CD401, is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD401 poly-A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (1) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (1) has antiarthritic, antirheumatic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (1) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.4%; Score 14.6; DB 1; Length 29; Best Local Similarity 69.0%; Pred. No. 2.4e+03; Matches 20; Conservative 0; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 29 BP; 22 A; 4 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3304 ATAGGATTTTTCTTTAGGAGATTTATTTT 3332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CD40L poly-A tract sequence SEQ ID NO:16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF74919 standard; DNA; 29 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-SEP-2000; 2000WO-US024966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-SEP-1999; 99US-0153625P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-MAY-2001 (first entry)
rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-244776/25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200119844-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1809
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XX
AC AAF7,
XX
AC AAF7,
XX
DT 23-M
XX
DX CD40
XX
Humal
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Humal
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HOMO
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The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (1) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAF74905 where A in the wild type sequence at position 331 (corresponding antirheumatic, immunosuppressive and antinflammatory activities, and can be used in gene therapy. (1) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L, is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly. A tract sequence which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as
                                                                                                                                                                                Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive; antiinflammatory; inflammatory disease; autoimmune disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.6; DB 1; Length 29; 59.0%; Pred. No. 2.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 29 BP; 22 A; 4 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3304 ATAGGATTTTTCTTTAGGAGATTTATTT 3332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                          CD40L poly-A tract sequence SEQ ID NO:20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Fig 3; 90pp; English.
                 AAF74923 standard; DNA; 29 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-2000; 2000WO-US024966.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF60442 standard; DNA; 39
                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 69.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein-RNA fusion; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-244776/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Li Y;
                                                                                                                                                                                                                                                                                                                    WO200119844-A1.
                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                               13-SEP-1999;
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                                                                                                     23-MAY-2001
                                                                                                                                                                                                                                                                                                                                                             22-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Crow MK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF60442;
                                                           AAF74923;
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Matches
                                         원
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated, purified nucleic acid, which is an altered CD40 ligand (CD40L) promoter (1) for CD40 ligand, having residues 331-455 of the sequence comprising 455 nucleotides given in AAP74905 where A in the wild type sequence at position 331 (corresponding to position -125) is replaced with C. (1) has antiarthritic, antirheumatic, immunosuppressive and antiinflammatory activities, and can be used in gene therapy. (1) is useful in the study, diagnosis and treatment of inflammatory and autoimmune diseases, as well as diseases in which elevated expression of CD40L is a factor, e.g., rheumatoid arthritis. The present sequence represents a CD40L poly-A tract sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New altered CD40L promoter for use in the study, diagnosis and treatment of a variety of inflammatory disorders and autoimmune diseases, such as rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; CD40L; promoter; CD40 ligand promoter; rheumatoid arthritis; diagnosis; antiarthritic; antirheumatic; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiinflammatory; inflammatory disease; autoimmune disease; ds.
                                                                             Length 29;
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0.4%; Score 14.6; DB 1; Length 29;
Best Local Similarity 69.0%; Pred. No. 2.4e+03;
Matches 20; Conservative 0; Mismatches 9; Indels
                                                                                                                           9; Indels
which is used in an example from the present invention
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                                       Sequence 29 BP; 22 A; 4 C; 0 G; 3 T; 0 U; 0 Other;
                                                                             0.4%; Score 14.6; DB 1;
69.0%; Pred. No. 2.4e+03;
tive 0; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NYRE-) NEW YORK SOC RELIEF RUPTURED & CRIPPLED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3304 ATAGGATTTTTCTTTAGGAGATTTATTT 3332
                                                                                                                                                                  3304 ATAGGATTTTTCTTTAGGAGATTTATTT 3332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD40L poly-A tract sequence SEQ ID NO:8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                      H.
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                                                                                                                                                                                                                                                                                                                      AAF74911 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                       23-MAY-2001 (first entry)
                                                                             Query Match
Best Local Similarity 69.0<sup>1</sup>
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-244776/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Li Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200119844-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-MAR-2001,
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AAF74911/

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Indels

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Location/Qualifiers

Unidentified

RESULT 1811

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(FERR/) FERRARI E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-NOV-2001
                  Zhong W,
                                                                                                                                                                                                                                                                                                  templates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stem_loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD27120
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                                                                                                                                                                                                                                                                             The present invention relates to a method for affixing a peptide acceptor to an RNA molecule through the formation of a covalent bond, noncovalent bond, or by chemical ligation. The method is useful for producing RNA-protein fusions which can be used for the isolation of proteins or mucleic aids with desired properties from large pools of partially or completely random amino acid or nucleic acid sequences. The present sequence is a DNA oligonucleotide used in the present invention
                                                                                                                                                                                                                                   an
                                                                                                                                                                                                              Affixing a peptide acceptor to an RNA molecule useful for producing fusion proteins for isolating proteins or nucleic acids with desired properties through attachment of a peptide acceptor to the 3' end of RNA molecule.
                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                           /note= "Pso-T, where Pso is a psoralen C2 amidite"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis C virus (HCV) NS5B replicase RNA synthesis template #1.
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                                                                                                                                                                                                                                                                                                                                                                     Score 14.6; DB 1; Length 39;
Pred. No. 2.88+03;
0; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis C virus; HCV; NS5B replicase; ss; RNA polymerase.
                                          /*teg= b
/mod_base= OTHER
/note= "C-Pu, where Pu is Puromycin-CPG"
                                                                                                                                                                                                                                                                                                                                                       Sequence 39 BP; 28 A; 5 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                             3262 TATTTTATTTGCTTTGTCCTTTTTCAGGA 3290
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                                                                                                                                                                                                                                                           Example 5; Page 19; 56pp; English.
                 /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK99271 standard; RNA; 40 BP
                                                                                                                    19-JUL-2000; 2000WO-US019653
                                                                                                                                       99US-0145834P
                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 69.0%;
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-APR-2000; 2000US-0195852P
                                                                                                                                                                           Lohse P, Wagner R;
        *tag≈ a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                             WPI; 2001-182803/18
                                                                                                                                                          (PHYL-) PHYLOS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ZHON/) ZHONG W. (HONG/) HONG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2002064771-A1
                                                                                 WO200107657-A1
modified base
                                   modified base
                                                                                                                                     27-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-OCT-2002
                                                                                                  01-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                             35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK99271;
                                                                                                                                                                           Kurz M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1813
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                                                                                                                                                        Novel replicase complex comprising hepatitis C virus NS5B replicase, a 3 nucleotide-long template to which a 2 nucleotide-long primer is annealed, and template and primer which do not form a stable duplex in the absence of HCV NS5B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of a reliable system for determining kinetic and thermodynamic constants of HCV NSSB-catalysed nucleotide incorporation and investigation of mechanistic inhibitors for mis-incorporation or chain termination. Specifically, the short RNA template and primer pairs are useful in screening assays which are used for determining kinetic, thermodynamic and mechanistic properties of NSSB replication and ultimately in the development of inhibitors of NSSB. Newly identified inhibitors of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 replicase activity may be used for developing anti-HCV pharmaceuticals. Sequences ABK99271-ABK99296 represent HCV NS5B replicase RNA synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA template with stable tetraloop, used to direct HCV RNA synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus; HCV replicase; non-structural protein 5B; NS5B;
lead compound; RNA polymerase; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9; Indels
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/bound_moiety= "Nucleotides 40-37"
/*tag= c
/bound_moiety= "Nucleotides 32-29"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40 BP; 29 A; 4 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3257 GAAGATATTTTATTTGCTTTGTCCTTTTT 3285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                              Example; Page 6; 17pp; English.
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Ferrari E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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9..32
                                                                       WPI; 2002-582330/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
Hong Z,
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The present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemloop at the 3' end capable of forming a copyback structure, a HCV non-structural protein 58 (NSSB), ATP, GTP, GTP, CTP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is and UTP nucleoside triphosphates (NTPS), where one of the NTP is and UTP nucleoside triphosphates (NTPS), where one of the NTP is and UTP nucleoside triphosphate for copy-back self-priming RNA synthesis of properties of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds or to characterise the biological relevance of lead compounds. The coptimal RNA templates can be used for developing a system to characterise HCV NSSB polymerase mechanistically and kinetically and for designing small RNA templates can be useful for developing a system to characterise. The nucleic acid synthesised by NSSB is detected by evaluating an entoradiograph of reaction products separated by gel electrophoresis. The present sequence is RNA template with a stable tetraloop used to direct RNA synthesis by RNA polymerase proceins of HCV, BVDV and polivirus. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                    Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.6; DB 1; Length 40; 59.0%; Pred. No. 2.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40 BP; 29 A; 4 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Fig 1B; 10pp; English.
                        99US-00309670
                                                                   99US-00309670
                                                                                                                                                                                                        Zhong W, Hong Z, Lau JYN;
                                                                                                                                                                                                                                                        WPI; 2002-096587/13
                                                                                                                                                            LAU JY N.
                                                                                                               ZHONG W.
                                                                 11-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                               (ZHON/)
                                                                                                                                                            (LAUJ/)
                                                                                                                                                                                                                                                                                                                                                                       buffer.
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The present invention describes an expressed sequence tag (EST), where the EST is an isolated, enriched, or purified nucleic acid sequence representing all or part of a gene, the expression of which, or its complementary sequence, in a cell identifies the cell as a developmentally competent or incompetent cell. Molecules which induce totipotence in one or more cells. Molecules which induced incompetence in a cell line are useful for inducing pregnancy in an animal and inhibiting totipotence. The molecules are also useful for treating a full term useful for treating a disease in an animal by inducing development of one or more cells of the animal into a specific cell type. The present sequence represents a bovine EST which is given in the exemplification of

An expressed sequence tag (EST), the expression of which, or its complementary sequence, in a cell identifies the cell as a developmentally competent or incompetent cell.

Example 16; Page 264; 584pp; English

Childs L;

Eilertsen KJ, Pfister-Genskow M,

WPI; 2002-351289/38.

06-JUN-2001; 2001US-00876143

(INFI-) INFIGEN INC

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Gaps

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Indels

0.4%; Score 14.6; DB 1; Length 41; 9.0%; Pred. No. 2.8e+03;

Query Match 0.4%; Score 14.6; D Best Local Similarity 69.0%; Pred. No. 2.8e Matches 20; Conservative 0; Mismatches

3306 AGGATTTTTTTAGGAGATTTATTTTT 3334

Sequence 41 BP; 3 A; 4 C; 6 G; 28 T; 0 U; 0 Other;

the present invention

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degenerative nervous system disease, graft versus host disease, hypersensitivity disease, infectious disease, neoplastic disease, acrophic gastritis; and interest disease, atrophic gastritis; and interest disease, atrophic gastritis; and in the solution nervous system disease, multiple sclerosis; Alzheimer's disease, hypersensitivity disease, type I hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; Infections disease, viral infection, type IV hypersensitivity; leprosy; infections disease, viral infection, and infection; schistosome;
                                                                                                                                                                                                                                                                                                                                                                                   filaria, bacterial infection, Mycobacterium, neoplastic disease;
lymphoproliferative disease; leukaemia, lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                            human; T-cell associated disease; Wheta; autoimmune disease;
 Human Vbeta gene repeat sequence #137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94US-00309335.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00263959
                                                                                    ADH70347 standard; DNA; 16
                                                                                                                                                           25-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         oreast cancer; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JS2002150891-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-MAR-1999;
                                                                                                                        ADH70347;
                                                       RESULT 1816
요
                                                                                          ö
                                                                                                                                            Gaps
                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                         Bovine embryonic germ (EG) cell cDNA EST 991115a CONTIG 45 #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Bovine; Bos taurus; EST; expressed sequence tag; totipotence;
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9; Indels

0; Mismatches

69.08;

20; Conservative

Matches

ઠે d

Best Local Similarity

3257 GAAGATATTTATTTGCCTTTGTCCTTTTT 3285

ABN73932 standard; cDNA; 41

03-JUL-2002 (first entry)

ABN73932;

development; gene; ss.

WO200194550-A2

Bos taurus.

13-DEC-2001

07-JUN-2001; 2001WO-US018576

07-JUN-2000; 2000US-0209874P

17-0CT-2002.

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descripted diseases which comprises a panel of nucleic acid primers associated diseases which comprises a panel of nucleic acid primers upocifically priming and allowing amplification of each Vbeta gene, vbetaRNA or colna. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases included diseases, concluding autoimmune disease, degenerative nervous system diseases, infectious diseases and neoplastic disease. Autoimmune diseases infolude Addison's disease, arobhic gastritis. Degenerative nervous system diseases include multiple of thypersensitivities and halpiner's disease. Hypersensitivity diseases include Type I hypersensitivities such as those present in doodpasture's syndrome and Type IV hypersensitivities such as those caused by armifested in leprosy. Infectious diseases include viral infections caused by the yeast genus Candida, parasitic infections such as those caused by whobbacterium. Neoplastic diseases include lymphoproliferative diseases such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                          autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  degenerative nearous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; arrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; Goodpasture's syndrome; type II hypersensitivity; Goodpasture's syndrome; type II hypersensitivity; leprosy; infectious disease; viral infection; HIV; fungal infection; parasitic infection; schistosome; filaria, bacterial infection, Mycobacterium, neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                 The invention relates to a kit for diagnosing and treating T-cell
                                                                                                                                                               Kit for diagnozing and treating T-cell associated diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.4%; Score 14.4; DB 1; Length 16; Best Local Similarity 93.8%; Pred. No. 1.5e+03; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; T-cell associated disease; Vbeta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 8 A; 1 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 541; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Vbeta gene repeat sequence #140.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3463 TATATATATCTATATA 3478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16 TATATATATATA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH70350 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2004 (first entry)
                                                                                                                WPI; 2004-059052/06
                                                                    Hood LE, Rowen L;
(HOOD/) HOOD L E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                breast cancer; ds
                      (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                      Vbeta gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADH70350;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADH70350
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US2002150891-A1

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The invention leaders to a ALL for diagnosing and treating Tection associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, ObetaMAN or CDNA. The Kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases.

Cr rejection and diagnosing and treating T-cell associated diseases including autoimmune diseases, diseases, infectious diseases, and neoplastic diseases. Autoimmune diseases include Addison's disease, arrophic gastritis. Degenerative nervous system diseases include Type carophic gastritis. Degenerative nervous system diseases include Type in hypersensitivities such as contact with allergens that lead to allergies. Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those caused by analises such as HIV, fungal infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by Wycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                       autoimmune, degenerative nervous system and infectious disease, comprises
nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a kit for diagnosing and treating T-cell
                                                                                                                                                                                                                                                    Kit for diagnozing and treating T-cell associated diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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0.4%; Score 14.4; DB 1; Length 16;
Best Local Similarity 93.8%; Pred. No. 1.5e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomere; budding yeast; eukaryotic; conserved region; phylogenetic relationship; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 7 A; 0 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 544; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S. cerevisae telomeric sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3465 TATATATCTATATA 3480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ51146 standard; DNA; 16 BP
                                                                   94US-00309335.
95US-00531241.
                                 99US-00263959.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 rararargrarara 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        [revised]
                                                                                                                                                                                                                 WPI; 2004-059052/06.
                                                                                                                                                                               Rowen L;
                                                                                                                         (HOOD/) HOOD L E.
                                                                                                                                          (ROWE/) ROWEN L.
                                 05-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     misc_feature
                                                                     19-SEP-1994;
                                                                                     19-SEP-1995;
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02-JUN-1994
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                                                                                                                                                                                                                                                                                                            Vbeta gene
                                                                                                                                                                             Hood LE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ51146;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1818
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Wright

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The present sequence represents a potential triple-helix forming region. It can be used to demonstrate the assay of the invention. The assay
                                                                                                                                                    Detection of eukaryotic pathogens, especially fungal or Plasmodium spp. by detecting telomerase activity.
                                                                                                                                                                                                                                                       The present sequence can be used in a novel method for detecting a eukaryotic pathogen in a patient. The method comprises obtaining a samp of somatic tissue or cells from the pathent, determining if telomerase activity is present and correlating this with the presence of the pathogen. The method is useful for diagnosis of fungal infections, especially a fungus of the genus Candida, Kluyveromyces, Saccharomyces, Sporothrix, Coccidioides, Histoplasma, Blastomyces, Paracoccidioides, Cryptococcus, Aspergillus, Mucor or Rhizopus, or malarial infections, especially Plasmodium vivax, P. ovale, P. malariae or P. falciparum. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Triple helix forming nucleotides 2771-2786 of Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Triple-helix forming region; Triplex formation; DNA detection; identification; bacteria; oncogene; virus; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 1.5e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 0 A; 0 C; 9 G; 7 T; 0 U; 0 Other;
                                                                             West MD,
              (UYCA-) UNIV CALIFORNIA SAN FRANCISCO.
(TEXA ) UNIV TEXAS SYSTEM.
                                                                             Mceachern MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Col 19-20; 168pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PROF-) PROFILE DIAGNOSTIC SCI INC.
                                                                                                                                                                                                                     Claim 5; Col 95-96; 82pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93US-00173489.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX14780 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 93.8'
Matches 15; Conservative
                                                                             Blackburn EH, Shay J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepburn AG, Wang C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-130384/11
                                                                                                                   WPI; 1998-041292/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX14780;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX14780/
셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from various budding yeast species. There is a great variety in the length and sequence complexity of these sequences compared to those of other eukaryotes they all have a 6 base conserved region at the 3' end. The telomeric relationships between these yeasts is fairly consistent with the phylogenetic relationship and it has been shown that C. tropicalis contains at least two forms of telomeric sequences. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                      Therapy and diagnosis of conditions involving telomerase using inhibitor - partic. for neoplasia, infection with pathogenic parasites and agerelated diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequences given in AAQ51138-46 represent telomeric sequences derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection; eukaryotic pathogen; telomeric nucleic acid sequence; telomerase activity; diagnosis; fungal infection; fungus; fungi; malaria! ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                     /*tag≈ b
/note= "May be truncated by multiples of 2 bp"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 0 A; 0 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                         West MD, Shay J, Wright W, Blackburn EH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fungal telomeric nucleic acid seguence.
"May be absent"
                                                                                                                                                                                                                                                                                                   UYCA-) UNIV CALIFORNIA SAN FRANCISCO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 30; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT96310 standard; DNA; 16 BP.
                                                                                                                                                                             93WO-US004546
                                                                                                                                                                                                                   92US-00882438
93US-00038766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 93.8%;
Matches 15; Conservative
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93US-00038766
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(first entry)
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  note=
                                   /*tag=
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1993-386602/48
                                                                                                                                                                                                                                                                                (GERO-) GERON CORP
                  misc_feature
                                                                                                                                                                             13-MAY-1993;
                                                                                                                                                                                                                     13-MAY-1992;
24-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-MAY-1992;
24-MAR-1993;
                                                                                              WO9323572-A1
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08-APR-1998
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two complementary strands of double stranded nucleic acid; (b) comparing the first value to a reference value (second reference value) of the second free energy parameter for the reference double stranded nucleic acid; and (c) if the test value and the second reference value of the second free energy parameter exhibit a test relationship that is the same as the preselected relationship, then preparing a first double stranded nucleic acid comprising all or part of the test nucleic acid, but if the test relationship is different than the preselected relationship, repeating step (a) and (b) on one or more additional test double stranded nucleic acids until an additional test double stranded nucleic acid is identified in which the test relationship is the same as the preselected relationship, and then preparing a first double stranded nucleic acid is comprising all or part of the additional test nucleic acid. AASB875-

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288882 represent the single stranded DNA molecules used to illustrate the method of the invention

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comprises adding a sample containing double-stranded DNA test sequences, e.g. containing the present sequence, to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (DV detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus
                                                                                                                                                                                          Gaps
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                                                                                                                                                               0.4%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 1.5e+03; tive 0; Mismatches 1; Indels
                                                                                                                                         Sequence 16 BP; 8 A; 0 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                Single stranded nucleic acid molecule (AT)2.
                                                                                                                                                                                                                                                                                                                                                                                                       Free energy parameter; thermodynamics; ss.
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                                                                                                                                                                                                                   923 TCTTCCTGTTCATCCT 938
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94US-00260200.
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Best Local Similarity 93.0.,
"..heg 15; Conservative
                                                                                                                                                                                                                                                                                                      AAZ88875 standard; DNA; 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Faldasz BD, Benight AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-APR-1994;
16-JUN-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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AAZ88875
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Gaps

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/ Match 0.4%; Score 14.4; DB 1; Length 16; Local Similarity 93.8%; Pred. No. 1.5e+03; hes 15; Conservative 0; Mismatches 1; Indels

Query Match

Matches

3464 ATATATATCTATATAT 3479

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1 ATATATAGCTATATAT 16

Sequence 16 BP; 7 A; 1 C; 1 G; 7 T; 0 U; 0 Other;

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This invention describes novel methods of producing double stranded nucleic acid molecules with a preselected value for a free energy parameter (e.g. a preselected T m or a preselected affinity for a nucleic acid binding ligand). The method involves preparing a first double stranded nucleic acid comprising a binding site for a nucleic acid binding ligand. The first value of a first free energy parameter of the first aciden uncleic acid has a preselected relationship with a first reference value of a first free energy parameter for a reference double-stranded nucleic acid comprising a reference binding site for the ligand. The first free energy parameter for a reference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Producing double stranded nucleic acid molecules with preselected values for free energy parameters such as affinity for nucleic acid binding
                                                                                                                         Single stranded nucleic acid molecule (AT) 2.
                                                                                                                                                        Free energy parameter; thermodynamics; ss.
                                                                                                                                                                                                                                                                                                                                                                             (UYNY ) UNIV NEW YORK STATE RES FOUND.
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94US-00224840.
94US-00260200.
                               AAZ88875 standard; DNA; 16 BP
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                                                                                           25-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                            Benight AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-194826/17.
                                                                                                                                                                                                                                                                                 11-DEC-1996;
                                                                                                                                                                                                                                                                                                                             08-APR-1994;
16-JUN-1994;
                                                                                                                                                                                                                                                                                                                17-JUN-1993;
                                                                                                                                                                                                                      US6027884-A.
                                                                                                                                                                                                                                                                                                                                                                                                            Faldasz BD,
                                                                                                                                                                                                                                                  22-FEB-2000
                                                                                                                                                                                       Synthetic.
                                                              AAZ88875;
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RESULT 1822
                  AAZ88875,
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This invention describes novel methods of producing double stranded nucleic acid molecules with a preselected value for a free energy parameter (e.g. a preselected Tm or a preselected affinity for a nucleic caid binding ligand). The method involves preparing a first double stranded nucleic acid comprising a binding site for a nucleic acid. binding ligand. The first value of a first free energy parameter of the first redenence value of a first free energy parameter for a reference caid comprising a reference binding with a first reference value of a first free energy parameter for a reference cauluse. The first free energy parameter for a reference cauluse is a characteristic of the binding of a ligand of incerest to its binding site and the preselected relationship is higher than, equal to or lower than (sic). The method comprises: (a) determining a test value for a test double stranded nucleic acid, comprising a test value for a test double stranded nucleic acid, comprising a test binding site for the ligand, of a second free energy parameter that is characteristic of the hybridization of the

Producing double stranded nucleic acid molecules with preselected values for free energy parameters such as affinity for nucleic acid binding

Lane MJ;

WPI; 2000-194826/17.

Disclosure; Col 69-70; 49pp; English.

ligands.

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binding of a ligand of interest to its binding site and the preselected relationship is higher than, equal to or lower than (sic). The method comprises: (a) determining a test value for a test double stranded nucleic acid, comprising a test binding site for the ligand, of a second free energy parameter that is characteristic of the hybridization of the two complementary strands of double stranded nucleic acid; (b) comparing the first value to a reference value (second reference value) of the second free energy parameter for the reference double stranded nucleic acid; and (c) if the test value and the second reference value of the second free energy parameter exhibit a test relationship that is the same as the preselected relationship, then preparing a first double stranded nucleic acid comprising all or part of the test nucleic acid, but if the cet relationship is different than the preselected relationship, the test double stranded nucleic acid is identified in which the test relationship is the same as the preselected relationship, and then preparing a first double stranded nucleic acid is identified in which then preparing a first double stranded nucleic acid comprising all or part of the test nucleic acid sating stranded nucleic acid seconds in the single stranded DNA molecules used to illustrate the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention provides a novel method for isolation of satellite sequences from genomic DNA that comprises fragmentation of the DNA by a method which is not dependent on base sequences, then selection of the satellite sequences from the obtained genomic library of high homogeneity. The method is useful for the isolation of microsatellite DNA sequences which can be used as DNA markers. The new method markedly. improves the efficiency of isolation of satellite sequences in comparison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolation of satellite sequences from genomic DNA for use as DNA markers comprises isolating a library with high homogeneity by DNA fragmentation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16 BP; 7 A; 1 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NORQ ) JAPAN MIN AGRIC FORESTRY & FISHERIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; Page 15; 35pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H. discus derived sequence #28
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                                                                                                                                                                                                                                                                                                                                                                                                                         method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-224692/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 independent cancer. The agents can be used to treat prostate, breast, cervical and ovarian cancers and to make hormonal drugs for retarding the conversion of a hormone-dependent cancer into a hormone-independent cancer. The drug can retard the change of hormone-dependent cancers into hormone-independent cancers effectively. The present sequence represents a PCR primer which is used in an example from the present invention
to prior art methods which are reliant on base sequences. Sequences AA298483-514 represent sequences from Haliotis discus, used in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a hormonal drug-containing agent (I) for retarding the conversion of a hormone-dependent cancer into a hormone-
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                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hormonal drug-containing agents for retarding conversion of hormone-dependent cancers, useful e.g. for treating prostate and breast cancers.
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Pred. No. 1.5e+03;
4; Mismatches 0; Indels
                                                                                          Length 16;
                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                            Human fibroblast growth factor antisense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 4 A; 5 C; 1 G; 2 T; 0 U; 4 Other;
                                                          Sequence 16 BP; 8 A; 6 C; 2 G; 0 T; 0 U; 0 Other
                                                                                     / Match 0.4%; Score 14.4; DB 1; Local Similarity 93.8%; Pred. No. 1.5e+03; nes 15; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 19; 31pp; Japanese.
                                                                                                                                                        2331
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                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                        2316 TCTGTGTGTGTGTGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matsutani E, Naito K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-303644/26.
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Best Local Similarity
                              of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200020034-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-OCT-1999;
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                                                                                                                                                                                                                                                                                                                               10-AUG-2000
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                                                                                                                                                                                                                                                                                                AAA29601;
                                                                                          Query Match
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Matches
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ID AAA2
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RESULT 1825

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The invention relates to detecting (MI) polymorphisms in a uridine diphosphate glucuronosyltransferase (UGT) gene promoter by determining the presence of five thymidine-adenine (TA) repeats in the promoter, where the presence of the five TA repeats correlates with increased expression of the gene. The method is used for detecting polymorphisms in cuseful for screening individuals for variation in glucuronidation activity, for optimising drug dosages for a patient, where the drugs cuseful for screening individuals for variation in glucuronidation activity of the drug dosages for a patient, where the drugs of e.g. Trinotecan or TAS-103) are glucuronidated by UGT (preferably UGTIAL) and the activity of the drug dosages for a patient, where the drugs of glucurodination. The method preferably involves obtaining DNA from an individual, amplifying all or part of a UGT gene promoter (UGTIAL) gene promoter. Consince the number of TA repeats of the drug amplified comprises all or part of UGTIAL promoter. The DNA and determining the number of TA repeats is determined amplified by a polymerase chain reaction and consisting of five TA repeats is determined by gel electrophoresis or by sequencing the amplified DNA. The polymorphism comprises an allele consisting of five TA repeats (TA)5, six TA repeats (TA)5, (TA)5,(TA)5, (TA)5,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8,(TA)8, (TA)8, (TA)8,(TA)8, (TA)8, (TA)8
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                                                                                                                                                                                                                                                                                                                                         Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, comprises determining the presence of five thymidine-adenine repeats in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ds; UGT1A1; promoter; Gilbert's syndrome; hyperbilirubinaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                with expression of the UGT gene, and the individuals sensitivity to xenobiotics is effected by glucuronidation activity. The methods preferably involve determining the presence of five, six or seven TA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.4; DB 1; Length 16; 33.8%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 16 BP; 8 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human UGT1a1 promoter polymorphism (TA)8 repeat.
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                                                                                                                                                                                                                                      Ratain MJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 7; Col 17; 13pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2823 TATATATACATATA 2838
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                                                                          99US-00251274.
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                                                                                                                                                                                                                                    Iyer L,
                                                                                                                                                                                  (ARCH-) ARCH DEV CORP
                                                                                                                                                                                                                                                                                          WPI; 2002-588597/63.
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Best Local Similarity
                                                                                                                                                                                                                                    Di Rienzo A,
                                                                          16-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                 the promoter
                                                                                                                              16-FEB-1999;
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                        28-MAY-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EXEXEXEX EXE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated human nuclear factor associated with dsRNA polypeptide useful for determining structure-function relationships and as affinity tag to identify and isolate interacting proteins that bind to the factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human nuclear factors associated with dsRNA (NFAR-1 and NFAR-2). These are transcriptional regulators and are thought to play a role in apoptosis and tumourigenesis. The coding sequence (found on chromosome 19p13.1-13.2) is useful as a probe to detect rearrangements in tumour cells and the protein is useful for determining structure-function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ds; UGI1A1; promoter; Gilbert's syndrome; hyperbilirubinaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    uridine diphosphate glucuronosyltransferase; Crigler-Najjar syndrome; UGT; polymorphism detection; TA repeat; glucuronidation; Irinotecan; TAS-103; xenobiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides the protein and coding sequences of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                           transcription regulator; chromosome 19p13.1-13.2; apoptosis; tumourigenesis; ds.
                                                                                                                                                                                                           Human; nuclear factor associated with dsRNA; NFAR-1; NFAR-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                       Human NFAR-1/NFAR-2 intron/exon junction sequence #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 16 BP; 3 A; 2 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human UGT1al promoter polymorphism (TA)8 repeat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Perkins DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 62; 73pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2673 GCCTCCCCACCTCCAG 2688
BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUN-2000; 2000WO-US015767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0138612P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCTCCCTACCTCCAG 1
AAF24305 standard; DNA; 16
                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Barber GN, Saunders L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BARB/) BARBER G N.
(SAUN/) SAUNDERS L.
(PERK/) PERKINS D J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-080688/09
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                                                                                                                                                                                                                                                                                                                                                                     WO200077205-A1
                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                     09-APR-2001
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                                              AAF24305,
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Gaps

; 0

1; Indels

RESULT 1826

Matches

à 셤 vivlemore401-10.rng

Human uridine diphosphate glucuronosyltransferase gene polymorphism #11.

16-JAN-2003 (first entry)

AAL50677;

Human; polymorphism; TA repeat; ds; UGT; thymidine-adenine repeat; uridine diphosphate glucuronosyltransferase gene promoter; UGT1A1; drug dosage optimisation; xenobiotic sensitivity.

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Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, comprises determining the presence of five thymidine-adenine repeats in
uridine diphosphate glucuronosyltransferase; Crigler-Najjar syndrome;
UGT; polymorphism detection; TA repeat; glucuronidation; Irinotecan;
TAS-103; xenobiotic.
                                                                                                                                                                     invention relates to detecting (M1) polymorphisms in a uridine
                                                                                                                                                                                                                                                                                                                                                                   Sequence 16 BP; 8 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                         Claim 7; Col 17; 13pp; English.
                                                                         99US-00251274
                                                              99US-00251274
                                                                                                  Iyer L,
                                                                                      (ARCH-) ARCH DEV CORP
                                                                                                              WPI; 2002-588597/63.
                          Homo sapiens
                                     US6395481-B1
                                                              16-FEB-1999;
                                                                          16-FEB-1999;
                                                                                                  Di Rienzo A,
                                                                                                                                            the promoter
                                                  28-MAY-2002
                                                                                                                                                                     The
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Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, involves determining number of thymidine-adenine repeats in the promoter.

Claim 7; Page 9; 13pp; English.

Ratain MJ;

Iyer L,

Rienzo AD,

WPI; 2002-740095/80.

99US-00251274

16-FEB-1999;

(ARCH-) ARCH DEV CORP

01-FEB-2002; 2002US-00061693

JS2002115097-A1

22-AUG-2002

Homo sapiens.

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The invention comprises a method for detecting polymorphisms in a uridine diphosphate glucuronosyltransferase (UGT) gene promoter (preferably UGTIAL). The method involves determining the number of thymidine-adenine (TA) repeats in the promoter - as the number of TA repeats correlates with expression of the UGT gene. The method of the invention is useful for detecting polymorphisms in a UGT gene promoter. The method of the invention is also useful in optimising drug dosages and predicting an individual's sensitivity to xenobiotics for drugs and xenobiotics that are glucuronidated by UGT. The present DNA sequence represents a UGT gene TA repeat polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human uridine diphosphate glucuronosyltransferase gene polymorphism #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; polymorphism; TA repeat; ds; UGT; thymidine-adenine repeat; uridine diphosphate glucuronosyltransferase gene promoter; UGT1A1; drug dosage optimisation; xenobiotic sensitivity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.4; DB 1; Length 16; 33.8%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 16 BP; 8 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2823 TATATATACATATA 2838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AALS0677 standard; DNA; 16 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TATATATATATA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JAN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 93.8
les 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAL50677;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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0.4%; Score 14.4; DB 1; Length 16; 13.8%; Pred. No. 1.5e+03; ve 0; Mismatches 1; Indels

93.8%;

Local Similarity 93.8 nes 15; Conservative

Query Match

2823 TATATATACATATA 2838

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AALS0677 standard; DNA; 16 BP.

RESULT 1828
AAL50677
ID AAL5067'
XX

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The invention describes a method use for treating increased rate of proliferation of a cell or extending the ability of a cell to replicate, or treating a disease associated with cell sensecence. The method comprises administering an agent to reduce loss of telomere length within the senescing cells. The method is useful for treating a condition associated with an increased rate of proliferation of a cell extending the ability of a cell to replicate, or for treating a condition associated with cell senescence e.g. neoplasia. A second method calsclosed in the invention is useful for treating a condition associated with an elevated level of telomerase activity within a cell e.g. cancer. Also disclosed is method useful for diagnosis of a condition associated with an increased rate of proliferation in a cell in an individual e.g. age-related macular degeneration, astrocytes associated with Alzheimer's disease and endothalial cells associated with atheroselerosis. This sequence represents a polymule of the study of telomere length
                                      Treating condition associated with cell senescence or increased rate of cell proliferation, by administering to cell an agent that derepresses telomerase in the senescing cells or that reduces loss of telomere
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nanocircle; telomere repeat sequence; cytostatic; ophthalmological; cancer; liver degeneration; macular; skin aging; gene therapy; biomedical research; tissue engineering; transplantation; ds; yeast.
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/note= "Each TG unit may be optionally absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 14.4; DB 1; Length 16; 3.8%; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomyces cerevisiae telomere repeat sequence DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and telomerase activity described in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 BP; 0 A; 0 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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"Optionally absent"
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                                                                                                                                                         Disclosure, Page 51; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC06894 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2318 TGTGTGTGTGTGTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae.
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7. .16
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WPI; 2003-066896/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1831
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                                                                                                                                                                                                                                        Detecting polymorphisms in uridine diphosphate glucuronosyltransferase gene promoter, useful for optimizing drug dosages for a patient, involves determining number of thymidine-adenine repeats in the promoter.
                                                                                                                                                                                                                                                                                                                                                                      The invention comprises a method for detecting polymorphisms in a uridine diphosphate glucuronosyltransferase (UGT) gene promoter (preferably UGT1A1). The method involves determining the number of thymidine-adenine (TA) repeats in the promoter — as the number of TA repeats correlates with expression of the UGT gene. The method of the invention is useful for detecting polymorphisms in a UGT gene promoter. The method of the invention is also useful in optimising drug dosages and predicting an individual's sensitivity to xenobiotics for drugs and xenobiotics that are glucuronidated by UGT. The present DNA sequence represents a UGT gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cell proliferation; cell senescence; telomere length; telomerase activity; cel replication; neoplasia; cancer; age-related macular degeneration; Alzheimer's disease; atherosclerosis; telomerase; telomerase inhibitor; immortalised cell; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomere length and/or telomerase activity related polynucleotide #57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 16 BP; 8 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blackburn EH;
                                                                                                                                                      Ratain MJ
                                                                                                                                                                                                                                                                                                                               Claim 7; Page 9; 13pp; English
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93US-00038766.
93US-00060952.
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                    01-FEB-2002; 2002US-00061693.
                                                               99US-00251274.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TA repeat polymorphism
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SHAY J.
WRIGHT W.
BLACKBURN E H.
                                                                                                           (ARCH-) ARCH DEV CORP
                                                                                                                                                      Rienzo AD, Iyer L,
                                                                                                                                                                                                  WPI; 2002-740095/80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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                                                               16-FEB-1999;
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24-MAR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABX50034;
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RESULT 1830

셤 ò

Best Loc Matches

(SHAY/) (WRIG/) (BLAC/) (WEST/)

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Gaps

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US2003190618-A1
                                  by the indexer
                                                                              Homo sapiens
                                                                                    09-OCT-2003
                                                                                                         Samal B,
                                         Query Match
                                                               AAD63066;
                                                                                               (SAMA/) (LIYY/) (HERM/) )
                                                                                                    (HOPP/) 1
(JOHE/)
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The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically primers amplification of each Vbeta gene, VbetaRNA or CNNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases including autoimmune diseases, degenerative nervous system diseases graft versus host diseases, hypersensitivity diseases, infectious diseases and neoplastic diseases. Autoimmune diseases include Addison's diseases atrophic gastritis. Degenerative nervous system diseases include multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; degenerative nervous system disease; areaft versus host disease; hypersensitivity disease; atrophic gastritie; disease; degenerative nervous system disease; multiple sclerosis; disease; hypersensitivity disease; type I hypersensitivity; dodases, type I hypersensitivity; allergy; type II hypersensitivity; dodasetuse; syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; fungal infection; candida; parasitic infection; schistosome;
tandem tag libraries of cDNAB. The step involves isolating a sample of mRNAB, amplifying the released tags, concatenating the amplified tags to form concatenated tags, amplifying and isolating the concatenated tags. The present sequence is human carboxypeptidase A3 tandem tag DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tilaria, bacterial infection; Mycobacterium; neoplastic disease;
lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 1.5e+03; tive 0; Mismatches 1; Indels
                                                                                                                       Sequence 16 BP; 0 A; 0 C; 7 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 540; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Vbeta gene repeat sequence #136.
                                                                                                                                                                                                                                                                       2330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94US-00309335.
95US-00531241.
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                                                                                                                                                                                                                                                                                                                     1 Grrrgrgrgrgrgr 16
                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH70346 Btandard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                       2315 GTCTGTGTGTGTGTGT
                                                                                                                                                                                                                          Conservative
                                                                                                                                                                    Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hood LE, Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast cancer; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2002150891-A1.
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19-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH70346;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                The invention relates to a novel nucleic acid nanocircle comprising at least two repeats of a telomere repeat sequence. The nanocircle of the invention demonstrates cytostatic and ophthalmological activities and may be useful during the diagnosis and treatment of cancer, liver therapy procedures. Purthermore, the nanocircle may be used to extend the lifespan of non-cancerous cell populations in culture, providing enhanced materials for biomedical research, tissue engineering and transplantation. The current sequence is that of the Saccharomyces cerevisiae telomere repeat sequence DNA of the invention. Note: this sequence is not displayed within the specification per se but was created
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Generating five prime biased tandem tag libraries of cDNAs by isolating a sample of mRNAs, amplifying the released tags, concatenating the amplified tags to form concatenated tags, amplified and isolating the
                                                                       Novel nucleic acid nanocircle comprising at least 2 repeats of a telomere repeat sequence, useful for extending length of telomere in vitro or in vivo, and for treating macular degeneration, and cancer in mammals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention discloses a method for generating five prime biased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tandem tag; concatenated tag; human; carboxypeptidase A3; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 14.4; DB 1; Length 16; 33.8%; Pred. No. 1.5e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16 BP; 0 A; 0 C; 9 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                         Disclosure; Page; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2318 TGTGTGTGTGTGTG 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAD63066 standard; DNA; 16 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Li Y, Hermida LC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 93.8
Matches 15; Conservative
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HOPPA N L.
JOHE K K.
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sclerosis and Alzheimer's disease. Hypersensitivity diseases include Type
                           I hypersensitivities such as contact with allergens that lead to allergies. Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by viruses such as HIV, fungal infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphopoliferative diseases such as leukaemias, Iymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16 BP; 8 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
88888888888888888
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0.4%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 1.5e+03; tive 0; Mismatches 1; Indels
                                                                2824 ATATATACATATAT 2839
                                                                                   15; Conservative
                Best Local Similarity
 Query Match
                                 Matches
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Gaps

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Human Vbeta gene repeat sequence #136.
    ADH70346 standard; DNA; 16 BP
             (first entry)
             25-MAR-2004
        ADH70346;
RESULT 1834
ADH70346/c
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degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; Alzhaimer's disease; hypersensitivity disease; type I hypersensitivity; doodpasture's syndrome; type II hypersensitivity; doodpasture's syndrome; type II hypersensitivity; leprosy; infectious disease; viral infection; thugal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer; human; T-cell associated disease; Vbeta; autoimmune disease; breast cancer; ds.

US2002150891-A1. Homo sapiens. 17-0CT-2002.

94US-00309335. 99US-00263959 05-MAR-1999; 19-SEP-1994; 19-SEP-1995;

Hood LE, Rowen L; (ROWE/) ROWEN L.

(HOOD/) HOOD L E.

Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a WPI; 2004-059052/06 /beta

Disclosure; SEQ ID NO 540; 164pp; English.

The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers

hormone dependent cancer that comprises a hormone group chemical agent and a differentiation inducing agent. Specifically, the hormone group chemical agent refers to an agonist derived from the luteinising hormone releasing hormone (LH-HH), whereas the differentiation inducing agent is preferably a fat soluble vitamin, or can be a nuclear receptor ligand, histone acetylation regulation drug or a DNA methylation regulation drug.

This invention relates to a novel therapeutic agent for the treatment of

Example 1; Page 14; 15pp; Japanese.

Novel therapeutic agent of hormone dependent cancer, comprising hormone group, chemical agent and differentiation inducing agent, useful for treating hormone dependent cancer in mammal.

WPI; 2004-113108/12.

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specifically priming and allowing amplification of each Vbeta gene,
VbetaRNA or CDNA. The kit is useful for diagnosing organ transplant
crejection and diagnosing and treating T-cell associated diseases
including autoimmune diseases, degenerative nervous system diseases,
crejections not diseases, hypersensitivity diseases, infectious diseases,
atrophic gastritis. Degenerative nervous system diseases,
atrophic gastritis. Degenerative nervous system diseases include multiple
colerosis and Alzheimer's disease. Hypersensitivity diseases include multiple
colerosis such as contact with allergens that lead to
leaflestes. Type II hypersensitivities such as those
and right in propersensitivities such as those
manifested in leprosy. Infectious diseases include viral infections
caused by viruses such as HIV, fungal infections such as those caused by
the yeast genus Candida, parasitic infections such as those caused by
whycobacterium. Neoplastic diseases include lymphoproliferative diseases
                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hormone dependent cancer; hormone; differentiation inducing agent; luteinising hormone-releasing hormone; LH-RH; fat soluble vitamin; prostate; ovarian; uterine; breast cancer; hormone independent cancer; anti-tumour; PCR; primer; s8; PDGF receptor.
                                                                                                                                                                                                                                                                                    such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Degenerate antisense PCR primer used to amplify the PDGF receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 1.5e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                           Seguence 16 BP; 8 A; 0 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  2824 ATATATACATATAT 2839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH23257 standard; DNA; 16 BP
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                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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Unidentified.
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WO9523225-A2
                    30-JAN-1995;
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03-APR-1997
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                                                                      Grimm S,
Modak A,
Tracz D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT53805;
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The present invention describes using the LH-RH agonist as a preventative or therapeutic agent for hormone dependent proteite, ovarian, uterine or breast cancer, and can also delay change in hormone independent cancer. Accordingly, the compositions of the invention are described as exhibiting high anti-tuneour activity. This oligonucleotide sequence is a degenerate PCR primer used to amplify the platlet-derived growth factor (PDGF) receptor, in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                    Bnzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                    Rat ICAM hammerhead ribozyme target sequence (nt. position 293)
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                                                                                                 . Match 0.4%; Score 14.4; DB 1; Length 16; Local Similarity 75.0%; Pred. No. 1.5e+03; les 12; Conservative 4; Mismatches 0; Indels
                                                                               Sequence 16 BP; 4 A; 5 C; 1 G; 2 T; 0 U; 4 Other;
                                                                                                                                          1798 AGTGACGTCTGGTCCT 1813
                                                                                                                                                                                                              AAT53487 standard; RNA; 17 BP.
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94US-00218934.
94US-00222795.
94US-00224483.
94US-00227958.
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94US-00271280.
94US-00291932.
94US-00291433.
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94US-00293520.
94US-00300000.
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94US-00311749.
94US-00314397.
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94US-00321993.
94US-00334847.
94US-00337608.
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15-APR-1994;
18-MAY-1994;
06-JUL-1994;
15-AUG-1994;
16-AUG-1994;
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23-SEP-1994;
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27-MAR-1997
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                                                                                                                                                                                                                     o DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW;
Karpeisky A, Kisich K, Matulic-Adamic J, Mcswiggen JA;
Pavco P, Bacjeleman L, Sullivan SM, Sweedler D, Thompson JD;
Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial; keumatoid arthritis; psoriasis; myocardial; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat ICAM hammerhead ribozyme target sequence (nt. position 2977)
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Pred. No. 1.6e+03;
4; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 5 A; 4 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 201; 407pp; English.
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94US-00357577.
94US-00363233.
95US-00380734.
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1 GAAGCUCUUCAAGCUG 16
                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 68.8°
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-351090/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                correct PI field.)
                                                                                                                                                                                                                                 Stinchcomb DT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-FEB-1995;
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The present sequence represents the preferred target sequence for an enzymatic nucleic acid, especially a harmerhead ribozyme, which cleaves the human c-myb sequence at the base position indicated in the descriptor line. The c-myb sequence was screened for optimal ribozyme target sites using a computer folding algorithm, and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes cleave the c-myb sequence and can be used to prevent smooth muscle cell hyperproliferation in restenosis, especially after coronary angioplasty,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New enzymatic nucleic acid molecules - cleave RNA produced by e.g. c-myb, for treating restenosis or cancer.
Human c-myb hammerhead ribozyme target sequence (nt. position 2894).
                                      Bnzymatic nucleic acid; hammerhead; ribozyme; cleavage; human;
smooth muscle cell; hyperproliferation; restenosis; cancer; c-myb;
coronary angioplasty; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse flk-1 VEGF receptor hammerhead ribozyme substrate #466.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stinchcomb DT, Draper K, Mcswiggen J, Jarvis T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 8 A; 0 C; 0 G; 0 T; 9 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 78; 128pp; English.
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95US-00373124.
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                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                18-MAY-1995;
                                                                                                                                                                                                                                                                                                                              18-MAY-1994;
13-JAN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Grimm S, Karpeisky A, Kisich K, Matulic-Adamic J, Mcswiggen JA;
Modak A, Pavco P, Beigleman L, Sullivan SM, Sweedler D, Thompson JD;
Tracz D, Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.
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18.8%; Pred. No. 1.6e+03;
ve 4; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; Page 204; 407pp; English.
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                   94US-00218934.
94US-0022795.
94US-00227958.
94US-00227958.
94US-00245736.
94US-00245736.
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94US-00291433.
94US-00292620.
94US-00293520.
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94US-00303039.
94US-00311486.
94US-00311749.
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94US-00316771.
94US-00319492.
94US-00321993.
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94US-00337608
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Best Local Similarity 68.8
Matches 11; Conservative
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                                                                                       15-APR-1994;
15-APR-1994;
18-MAY-1994;
06-JUL-1994;
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16-AUG-1994;
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19-AUG-1994;
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04-NOV-1994;
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                                                                                                                                                                                                      The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (WEGF). A patient (preferably human) having a condition associated with the level of the fins-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, coular diseases, psoriasis and rhematoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                         Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Parallel automated synthesis of oligo:nucleotide(s) on alkylamino-modified support - particularly for producing oligomer chips for use in hybridisation experiments or as source of high quality bio:polymer(s).
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                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.4; DB 1; Length 17; 31.2%; Pred. No. 1.6e+03; ve 2; Mismatches 1; Indels
                                                                                               Escobedo J;
                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 5 A; 3 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Plasmid pTZ18R derived oligomer chip #9
                                                                                               Stinchcomb D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM
                                                                                                                                                                                    Claim 4; Page 137; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                1608 GAAGTGCATCCACAGG 1623
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           96WO-US017480.
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                               95US-0005974P
                                           96US-00584040
                                                                                                                                                                                                                                                                                                                                                               81.2%;
                                                               (RIBO-) RIBOZYME PHARM INC. (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV09832 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                Local Similarity 81.2
Les 13; Conservative
                                                                                               P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Weiler J, Höheisel J;
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                                                                                                                      WPI; 1997-259017/23,
           25-0CT-1996;
                                26-OCT-1995;
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                                           11-JAN-1996;
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                                                   Oligonucleotides AAV09824-V09844 are used to explain a novel method for the parallel automated synthesis of oligonucleotides on an alkylaminomodified matrix surface. This method involves applying 3-succinate derivatives of protected nucleosides to the matrix surface. The oligomer chips produced this way are used in hybridisation experiments and as source of high quality biopolymers (for polymerase chain reaction, sequencing, or as probes). The method allows the simple production of a wide range of different oligonucleotides
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                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 14.4; DB 1; Length 17; 33.8%; Pred. No. 1.6e+03; ve 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 2 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
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Example, Fig 5C; 26pp; German.
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ses 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-077103/07.
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The capulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule opssessing an NCH mocif), a G-cleaver (cleaving RNA with an NGN with an NGN triplet), a zinzyme (cleaving RNA with an NGN triplet), a zinzyme (cleave RNA contacted of a divalent cation that is preferably MG<sup>2</sup>+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level conforts. The reatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-creat lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic lymphocytic lymphoma, (MLL), immunocytoma (MCL), small B-cell lymphocytic lymphoma, immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunocytoma (MCL), small B-cell lymphocytic lymphoma, immunocytoma (M
                                                                                                                                                                                                                                                                                                                                                             Human; se; antisense therapy, cytostatic; antiinflammatory; haemostatic; cerebroprotective; notropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-dleaver; amberzyme; zinzyme; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer; s disease; miltiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 30; Page 152; 200pp; English.
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                                                                                                                                          ABK03424 standard; RNA; 17 BP
2 CTTGCATGCCTGCAGG 17
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06-MAR-2000; 2000US-0187128P
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                                                                                                                                                                                                                                                                                                              Human CD20 G-cleaver #39.
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MCSWIGGEN J.
CHOWRIRA B M.
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                                                                                  RESULT 1842
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presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contracted with a cell to reduce NoGo activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CVA, stroke). Alzhaimer's disease, dementia, multiple sclerosis (MS), parkinson's disease, taxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence is a G-cleaver molecule of the invention
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Synthetic.
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(MCSW/)
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Matches
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Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic lymphoma, leukaemia, HV (human immunodeficiency virus) associated NHL, lymphoma, immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCL), immunocytoma (IMC) small B-cell lymphocytic lymphoma, immunodeficiency virus) associated NHL, lymphoma, immunodeficienc
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states which respond to the modulation of NOGO expression. The present
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                               88; Page 98; 200pp; English.
 nervous system injury.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1844
                               Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be
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                                                                                                                             New nucleic acid(s) for regulating the Grb2-related with Insert Domain
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   Ellis JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ellis JH;
                                                                                                                                                               (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
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Mcswiggen JA, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; Grb2-related with Insert Domain; GRID; T-cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human GRID Amberzyme substrate oligonucleotide #147.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 3 A; 8 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Von Carlowitz I, Mcswiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
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                                                                                                                                                                                                                                                              Claim 4; Page 66; 108pp; English.
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   Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GCCCCAGCAGCUGCAG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 87.5%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          leukaemia, cytostatic, ss.
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                                                                WPI; 2001-550088/61.
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   Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL47247;
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AAH91733 standard; DNA; 17
                                     AAH91733;
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              was
                                                                                            Gaps
           chemotherapy and cyclosporin treatment. The present oligonucleotide used to illustrate the invention
administered in conjunction with other therapies such as radiation
                                                                                                                                                                                                                                                                                  Human; Grb2-related with Insert Domain; GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ellis JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid(s) for regulating the Grb2-related with Insert I (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
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                                                                   0.4%; Score 14.4; DB 1; Length 17; 75.0%; Pred. No. 1.6e+03; Live 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.4; DB 1; Length 17; 75.0%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Von Carlowitz I, Mcswiggen JA, Hamblin PA,
                                             Sequence 17 BP; 2 A; 2 C; 9 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                 Human GRID zinzyme substrate oligonucleotide #96.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 73; 108pp; English.
                                                                                                               2006 TGGTGGAGGACCTGGA 2021
                                                                                                                                                                                              ВР
                                                                                                                             2 UGGUGGAGGUCCUGGA 17
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                                                                                                                                                                                              ABL47012 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                          (first entry)
                                                              Query Match
Best Local Similarity 75.09
Matches 12, Conservative
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                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                     ABL47012;
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The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colities and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                       Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis;
                                                                                                                                   single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                     Human inflammatory bowel disease associated polymorphic site #808.
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                                                                                                                                                                                                                                                                                                                      /*tag= a
/note= "SNP, optionally T or A at this position"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WHED ) WHITEHEAD INST BIOMEDICAL RES. (BLLI-) BLLIPSIS BIOTHERAPEUTICS CORP.
                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1165 AAATGGGAGCTGTCTCG 1181
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ID ABN08005 standard; DNA; 17
(first entry)
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misc_feature
                                                                                                                                                                                                                    Homo sapiens.
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  09-OCT-2001
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Gaps

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1; Indels

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RESULT 1847

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cacids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for cacressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunospens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describing in onisation, as the rapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration and in vaccines or for replacement therapy. The concentration and skeletal muscle disorders hGDMLP-1 is localised to chromosome 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hanzel DK, Rank DR, Chen W, Shannon ME;
skeletal muscle disorder; amplicon; screening; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 6 A; 5 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 7997; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                        30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
                                                                                                                                                                                            2000US-0237456P.
2000US-0234687P.
2000US-0236359P.
                                                                                                                                                         25-MAY-2001; 2001WO-US016981
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2001WO-US000670
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AEOM-) AEOMICA INC
                                                                           WO200192524-A2
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                                                                                                                                                                                                                                                                                                                                                                          30-JAN-2001;
30-JAN-2001;
                                      Homo sapiens
                                                                                                                                                                                                                   21-SEP-2000;
27-SEP-2000;
                                                                                                                  06-DEC-2001
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Gaps ö 0.4%; Score 14.4; DB 1; Length 17; 33.8%; Pred. No. 1.6e+03; ve 0; Mismatches 1; Indel8 Query Match

Best Local Similarity 93.8%;
Matches 15; Conservative

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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                               Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7995.
                                                                                                                                                                                                                                                                                  Shannon ME;
                                                                                                                                                                                                                                                                                  Chen W,
                                                                                                                                                                                                                                                                                  Rank DR,
                                                                                                                                                                                                                                                                                  Hanzel DK,
      BP.
                                                                                                                                    25-MAY-2001; 2001WO-US016981.
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     ABN08003 standard; DNA; 17
                                  (first entry)
                                                                                                                                                                                                                                                                                  Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                     (AEOM-) AEOMICA INC.
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                                                                                                         WO200192524-A2.
                                                                                           Homo sapiens
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                                 29-MAY-2002
                                                                                                                      06-DEC-2001,
                    ABN08003;
ABN08003
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 7995; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunospens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP-proteins, as specific biomolecule and/or amount specifically of hGDMLP-proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therefore a section of hGDMLP-1 may be used for diagnosing a conduction, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and skeletal muscle disorders, hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1 in particular heart and skeletal muscle disorders, hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the CDMLP-1 sequence of this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fire, who int/pub/published_pct_sequence

Sequence 17 BP; 6 A; 6 C; 3 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polypeptide, for raising antibodies that recognize hGDWLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDWLP-1.
                                                                                                                                                                                                                                                                                                                                Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
skeletal muscle disorder; amplicon; screening; ss.
                                      Gaps
                                                                                                                                                                                                                                                                                                  Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:2002
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0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03; ive 0; Mismatches 1; Indels
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27-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234687P.
04-OCT-2000; 200US-0236359P.
04-OCT-2000; 200US-00206513.
30-JAN-2001; 200UMO-US000667.
30-JAN-2001; 200UMO-US0006665.
30-JAN-2001; 200UMO-US0006666.
30-JAN-2001; 200UMO-US000666.
30-JAN-2001; 200UMO-US000666.
30-JAN-2001; 200UMO-US000667.
30-JAN-2001; 200UMO-US000667.
30-JAN-2001; 200UMO-US000667.
                                                                  1992 CACCTTCAAGCAGCTG 2007
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                Local Similarity 93.8
1es 15; Conservative
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Query Match
Best Local S:
Matches 15
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production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence date for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
skeletal muscle disorder; amplicon; screening; ss.
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30-JAN-2001; 2001WO-US000663.
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30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
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Best Local Similarity 93.88
Matches 15; Conservative
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nucleic acids can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify composed in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule computer probes for surface-enhanced laser desorption ionisation, as therefore that specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a disorder sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO cat fib. wipo.int/pub/published_pct_sequence
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21-SEP-2000; 2000US-0234687P.
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as steadards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The collisor desequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart can disbetatal muscle disorders. hGDMLP-1 is localised to chromsome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence and the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed formed the present invention.
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                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                              Disclosure; SEQ ID NO 2005; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1487 GCCCCCCGGGCCTGGA 1502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-SEP-2000; 2000US-0236359F.
04-OCT-2000; 2000GB-00024263.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN02014 standard; DNA; 17 BP
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                 WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAY-2000;
21-SEP-2000;
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ABN02014/c
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30-JAN-2001; 2001WO-US000667. 30-JAN-2001; 2001WO-US000668. 30-JAN-2001; 2001WO-US000669. 30-JAN-2001; 2001WO-US000670. 23-MAX-2001; 2001US-00864761.

Shannon ME;

Chen W,

WPI; 2002-684061/74.

(AEOM-) AEOMICA INC.

Shannon M;

2001WO-US000665

30-JAN-2001; 30-JAN-2001;

28-JAN-2002; 2002EP-00001165.

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 (an be used in gene therapy and vaccine production. The hGDMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 mucleic acids in samples, as amplification substrates, to compare a simmunospens to raise antibodies that specifically recognise hGDMLP-1 proteins or polypeptides may be consed as immunospens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically of hGDMLP-1 proteins, as specific biomolecule and/or amount specifically of hGDMLP-1 proteins, as specific biomolecule captures probes for surface-enhanced laser desorption ioniation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The conduction, and in the exemplification of HGDMLP-1 may be used for diagnosing a disorder associated with the expression of HGDMLP-1 caquence in the exemplification of the present sequence of this patent an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO capture for this patent did not form part of the printed capture for this patent did not form part of the printed capture for this patent did not form part of the printed capture for this patent did not form part of the printed capture for this patent did not form part of the printed capture for this patent did not form part of the printed captures.
                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                                                                         Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 2006; 214pp; English.
                                                                                                                                                                                                                                                                                            Hanzel DK,
                       30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
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                                                                                                                                                                                                                                                                                         Ji Y, Penn SG,
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30-JAN-2001;
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Gaps ; 0 Ouery Match 0.4%; Score 14.4; DB 1; Length 17; Best Local Similarity 93.8%; Pred. No. 1.6e+03; 1; Indels 0; Mismatches

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> Human POSHL1 scanning oligonucleotide SEQ ID NO 441. ABV89728 standard, DNA, 17 BP 23-DEC-2002 (first entry) ABV89728; RESULT 1854
> ABV89728
> XX
> AC ABV89728
> AC ABV89728
> AC ABV89726
> XX
> DT 23-DEC-2
> XX
> C Human PC
> XX
> KW Human, EW
> XW Grpe
> KW Gr

Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.

EP1239051-A2

Homo sapiens

Homo sapiens

11-SEP-2002.

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBA399), a sequence having 65% sequence of 60min, caids (SI, ABBA399), a sequence having 55% dentity to (SI), (SI), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Thuman POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTBases as well as downstream components of the signal transduction pathway. (I) is useful and contentifying a specific binding partner. (I) and nucleic acids (II) concoding (I) are useful for diagnosing, monitoring disease and treating canser, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating care useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomorleotide useful in examples of the invention. Note: The present sequence information supplied to printed specification, but is based on sequence information supplied to
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Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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                                                                                                                                             Example 2; SEQ ID NO 441; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Derwent by the Buropean Patent Office
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les 15; Conservative
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Matches
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EP1239051-A2

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Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated SH3 domain (POSH)-like signalling acids (SI, ABB8399), a sequence (II), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence identity to (SI), (SI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene-product that functions as adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying aspecific binding partner. (I) and nucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 443; 60pp + Sequence Listing; English.
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30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
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2001WO-US000666
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Best Local Similarity 93.84
Matches 15; Conservative
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                             Homo sapiens.
                                                                           EP1239051-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (61, ABB8399), a sequence having 65% sequence identity to (81), (81) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (1) and nucleic acids (II) encoding (I) are useful for idagnosing, monitoring disease and treating caused by aleared expression of human POSHL1 including disgnosing and treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomoucleotide useful in examples of the invention. Note: The present sequence information supplied to printed specification, but is based on sequence information supplied to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated SH3 domain (POSH)-like signalling
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30-JAN-2001; 2001WO-US000665
30-JAN-2001; 2001WO-US000666
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30-JAN-2001; 2001WO-US000668
30-JAN-2001; 2001WO-US000669
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23-MAY-2001; 2001US-00864761.
10-OCT-2001; 2001US-0328205P.
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                                                11-SEP-2002
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Gaps . 0

1; Indels

Human POSHL1 scanning oligonucleotide SEQ ID NO 445.

Human, POSHL 1; SH3 domain, POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.

RESULT 1856

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ABV89730
ID ABV8
XX ABV8
XX ABV8
XX ABV8
XX ABV8
XX Huma
XX Huma
XX Huma
XX Huma

virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;

Amberzyme; Zinzyme; ss

West Nile Virus WO200268637-A2. 36-SEP-2002 19-OCT-2001; 2001WO-US048350. 20-OCT-2000; 2000US-0242411P.

(RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L.

(MCSW/) MCSWIGGEN J A.

Mcswiggen JA;

Blatt L,

WPI; 2002-706994/76.

WNV minus strand Amberzyme substrate SEQ ID NO 14913.

Nile Virus; antiinflammatory;

cytostatic; hepatotropic;

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3999), a sequence having 65% sequence of 6730 amino acids (SI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for independent of vaccines and in secured by altered expression of human POSHL1 including disease and treating caused by altered expression of human POSHL1 including diagnosing and creating caused by altered expression of human POSHL1 including diagnosing and creating caused, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating care useful for measuring and for surveying gene expression and creating care useful for measuring and for surveying gene expression and creating care useful for measuring and for surveying gene expression and creating present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence information supplied to become
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL -1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; SEQ ID NO 445; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 2 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Derwent by the European Patent Office
                                                                                                                                                                                                                                                                         30-JAN-2001; 2001WO-US006664.
30-JAN-2001; 2001WO-US006665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001US-00864761.
                                          gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     549 GCTGCCGGCCAACCAG 564
                                                                                                                                                                                                               28-JAN-2002; 2002EP-00001165
                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-OCT-2001; 2001US-0328205P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-684061/74
                                                                                                                                                                                                                                                         30-JAN-2001;
30-JAN-2001;
                                                                                    Homo sapiens
                                                                                                                          EP1239051-A2
                                                                                                                                                                       11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shannon M;
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New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.

Claim 23; SEQ ID NO 14913; 495pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WNV Hammerhead Ribozyme substrate SEQ ID NO 532.
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BP.

ACN14910 standard, RNA; 17

RESULT 1858 ACN14910/c 22-APR-2004 (first entry)

ACN14910;

UXXXE

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is concoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p1.2 and MDZ12 is encoded at chromosome 16p2.2 cor in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3. MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids and also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapoutic agents for gene therapy or as the present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD21; MD21; MD21; chromosome 7q22.1; chromosome 6p21.3-22.3; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                     New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 7 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human MDZ3 scanning oligonucleotide SEQ ID 129.
                                                                                                                                                                                                                                                                                                                                                                                Example 8; SEQ ID NO 130; 103pp; English.
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  30-JUL-2002; 2002EP-00016874.
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                                                  02-AUG-2001; 2001US-00922181
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                                                                                                                                                   Shannon M, Gu Y, Nguyen C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                    (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, invocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DMAZyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nuclectides and a 3' end andification of a 3' -3' inverted abasic molety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid molecule that modulates replication of West Nile Virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 23; SEQ ID NO 532; 495pp; English.
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                                                                                                                                                                                                                                                     20-OCT-2000; 2000US-0242411P
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                                                                                                                                                                                                                                                                                                                                                                                                           Blatt L, Mcswiggen JA;
Amberzyme; Zinzyme; ss.
                                                                                                                                                                                                                                                                                                                              (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-706994/76.
                                               West Nile Virus.
                                                                                                    WO200268637-A2.
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 7422.1, MDZ4 is encoded at chromosome 7422.1, MDZ7 is encoded at chromosome 1621.3-22.2, MDZ7 is encoded at chromosome 1621.3 encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12, The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer to detect and characterize gross acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 7 A; 2 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                   Example 8; SEQ ID NO 129; 103pp; English.
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WPI; 2003-423107/40.
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0.4%; Score 14.4; DB 1; Length 17; 33.8%; Pred. No. 1.6e+03; ve 0; Mismatches 1; Indels 1295 TGAAGATGCTGAAAGA 1310

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Gaps

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(first entry) 21-MAR-2003 ABZ61917; RESULT 1862
ABZ61917/c
AC ABZ6191
XX
XX
DT 21-MAR-;
XX
DY Human H
XX
Human;
XX
CO HOMO Sa
C

Human H-Ras DNAzyme target #708.

Human, ribozyme, short interfering RNA, siRNA, HER2, K-Ras, enzymatic nucleic acid, H-Ras, N-Ras, HIV, cytostatic, anti-HIV, anti-rheumatic, cancer, AIDS, ss.

Homo sapiens.

WO200297114-A2.

05-DEC-2002

29-MAY-2002; 2002WO-US016840.

29-MAY-2001; 2001US-0294140P. 06-JUN-2001; 2001US-0296249P. 10-SEP-2001; 2001US-0318471P.

(RIBO-) RIBOZYME PHARM INC.

Mcswiggen J;

WPI; 2003-140484/13.

Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.

Claim 58; Page 124; 185pp; English.

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                                 acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immundeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing also useful for reducing breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ66531, ABZ66520 - ABZ66524, ABZ66531, ABZ66530 - ABZ66585 represent substrate/target sequences for the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or panoreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ65531, ABZ65521, ABZ65524,
                    novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03; Ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 9 C; 3 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 58; Page 114; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human H-Ras DNAzyme target #164.
                                                                                                                                                                                                                                                                                                                                                                                                2900 CAGGAGGCAGGCATGG 2915
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                      Local Similarity 93.8%;
nes 15; Conservative
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                    invention relates to a
                                                                                                                                                                                                                                       ribozymes of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ61373;
                                                                                                                                                                                                                                                                                                                       Query Match
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Matches
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3626 GGGCCCTGAGTCTGGG 3641
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08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
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                                                                                                                                                                                                                                                                                                                            16 GGGCCCTGACTCTGGG 1
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                                                                                                                                                                                                                                      Query Match
Best Local Similarity 93.8
Matches 15; Conservative
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Roberts E;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Macejak
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAVCO P.
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Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                ACD53091;
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(BLAT/)
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(ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PAVC/)
(LEEP/)
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                                                                                                                                                                                                                                                                                                                                                                     RESULT 1865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNN. The nucleic acid molecules include antisense and enzymatic nucleic acids as hammerhead ribozymes, DNAzymes, inczymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                               RNA stability; RNA expression; RNA synthesis; antisense; enzymetic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; decoyerse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
ABZ66530 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention
                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecule, Hepatitis C virus, HCV, Hepatitis B virus, HBV;
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                                                                         0.4%; Score 14.4; DB 1; Length 17; 33.8%; Pred. No. 1.6e+03; ve 0; Mismatches 1; Indels
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                                              Sequence 17 BP; 0 A; 3 C; 12 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Morrissey D,
                                                                                                                                                                                                                                                                                                                           HBV hammerhead ribozyme substrate sequence #8.
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                                                                                                                                    2158 CCCCGGCCCCACCCA 2173
                                                                                                                                                                                                                                        ACD50343 standard; RNA; 17 BP
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08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2001; 2001US-00817879
                                                                                         93.88;
                                                                                                                                                   16 CGCCGGCCCCACCCA 1
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BLATT L.
                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                        15, Conservative
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Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
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                                                                                          Best Local Similarity
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ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis B virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200281494-A1.
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                                                                                                                                                                                                                                                                                                23-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection.
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                                                                            Query Match
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(BLAT/)
(MACE/)
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(MORR/)
(PAVC/)
(LEEP/)
(DRAP/)
                                                                                                                                                                                                        RESULT 1864
ACD50343/c
                                                                                                        Matches
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are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HBV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amberzyme, G-cleaver ribozyme, decoy molecule, aptamer, they reverse transcriptse; Enhancer I region; viral replication; degenerative, disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                            ribozyme, inozyme, G-cleaver, zinzyme, DNAzyme or amberzyme sequences disclosed in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 3 A; 7 C; 5 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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The present invention relates to nucleic acid molecules which modulate
the synthesis, expression and/or stability of Hepatitis C virus (HCV) or
Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense
and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes,
inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed
are nucleic acid decoy molecules and aptamers that bind to HBV reverse
transcriptase and/or HBV reverse transcriptase primer sequences, as well
as oligonuclectides that specifically bind the Enhancer I region of HBV
DNA. The nucleic acids may be used to modulate the expression of HBV
compounds and/or potential therapies directed against HBV, and compounds
compounds and/or potential therapies directed against HBV, and compounds
compounds of the invention are useful for the treatment of degenerative and
disease states related to HBV and HCV infection, replication and gene
expression such as cirrhosis, liver failure, and hepatocellular
carcinoma. The present sequence represents a substrate for one of the HBV
ribozyme, inozyme, G-cleaver, zinzyme, DNAzyme or amberzyme sequences
                  hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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O
Novel compound useful for treating cirrhosis, liver failure,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 14.4; DB 1; Length 17; 13.8%; Pred. No. 1.6e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 2 A; 10 C; 2 G; 0 T; 3 U; 0 Other;
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                                                                              Example 1; Page 164; 387pp; English.
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08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
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BLATT L.
MACEJAK D.
MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             853 GAGGAGGAGCTGGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.4
Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200281494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-0CT-2002
                                          infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACD54765;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) 1
(BLAT/) 1
(MACE/) 1
(MCSW/) 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1866
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inczymes, zinzymes, amberzymes, and G-cleaver ribozymes. DNAzymes, compounds and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV DNA. The nucleic acids may be used to modulate the expression of HBV C genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HBV intoxyme, inoxyme, G-cleaver, zinzyme, DNAzyme or amberzyme sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis; virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
                                                                                                                                                                                      Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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П
                                                                                                     Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.4; DB 1; Length 17; Pred. No. 1.6e+03;
                                                                                                    Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumour suppression/reversion associated nucleotide #2441.
                                                                                                    Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A; 4 C; 3 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.4%; Score 14.4; best Local Similarity 62.5%; Pred. No. 1.6e
                                                                                                                                                                                                                                                          Example 1; Page 188; 387pp; English
                                                                                                    Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disclosed in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2776 TICCGGAAACTAGIGT 2791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB42118 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-2002; 2002WO-IB004219.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-2001; 2001FR-00011981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 UUCCGGAAACUACUGU 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
                                                                                                    Blatt L, Macejak D,
Draper K, Roberts E;
MORRISSEY D.
                                                                                                                                                    WPI; 2003-229207/22.
                                                                ROBERTS E.
                                LEE P.
DRAPER K.
                   AVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003040369-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis.
                                                                                                                                                                                                                           infection.
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                                                                   ROBE/)
MORR/)
                                 LEEP/)
                PAVC/)
                                                DRAP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1867
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ADI47758;
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                                                                                                                                                                                                                                                                                                                                                                         also be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
ઠે
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                                                                                                                                                                      fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying udantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as recombinant models. The nucleotides (also vectors containing the mand cells containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment of tumours of viral infections or diseases characterized by development of tumours cor cell degeneration (e.g. Alzheimer's disease or schizophrenia).

Analysis of the expression of the nucleotides and polypeptides can and or prognosis of these diseases. The nucleotides and polypeptides can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis; virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
                                                                           New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                  also be used to screen for their specific interactive molecules, potentially useful for treating diseases associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 2 A; 5 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                Disclosure; Page 317; 771pp; French.
                             Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1299 GATGCTGAAAGACGAT 1314
(MOLE-) MOLECULAR ENGINES LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression of the nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB45815 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15; Conservative
                             relerman A, Amson R,
                                                        WPI; 2003-441574/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003040369-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-MAY-2003,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1868
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ADB45815/
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Tuijnder M;

Amson R,

relerman A,

(MOLE-) MOLECULAR ENGINES LAB

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free invention relates to the isolation of 6327 nucleotide sequences, fragments of at least 18 consecutive nucleotides of these mucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the the nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tumour cuppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing them and cappression or the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours or call degeneration (e.g. Alzheimer's disease or schizophrenia).

Analysis of the expression of the nucleotides can be used for disagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can and/or prognosis of these diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour suppression; tumour reversion; apoptosis; virus resistance; cytostatic; virucide; neuroprotective; nootropic; neuroleptic; probe; primer; PCR; gene chip; antisense; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ds; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used to screen for their specific interactive molecules, ally useful for treating diseases associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human tumour suppression/reversion-related DNA sequence SeqID261.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 5 A; 2 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                   Disclosure; Page 749; 771pp; French.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2380 CATCTTGCCTCCAGGT 2395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression of the nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI47758/c
ID ADI47758 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.88;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 93.8
hes 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              relerman A, Amson R,
WPI; 2003-441574/41.
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Gaps

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Indels

Pred. No. 1.6e+03;); Mismatches 1;

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15; Conservative

Best Local Similarity Matches 15, Conserv

93.8%;

2110 AGCTCCAGCTCCTCAG 2125

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17 AGCTCCAGCTTCTCAG 2

0.4%; Score 14.4; Query Match

Sequence 17 BP; 4 A; 3 C; 6 G; 4 T; 0 U; 0 Other;

DB 1; Length 17;

NAC-1 protein. The DNA and protein sequences of the invention are useful for treating or preventing a disorder associated with decreased or increased expression or activity of human NAC-1. The DNA and protein sequences are also useful for treating drug addiction and cancer. The present DNA sequence represents a probe for the human NAC-1 gene.

The invention comprises the amino acid and coding sequences of the human

Example 2; SEQ ID NO 166; 96pp; English

0.4%; Score 14.4; DB 1; Length 17;

Query Match

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in the phenomena of the control of compounds with a cytostatic, virucide, neuroprotective, contropic or neuroleptic activity. The DNA sequences may be useful for the nootropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one compount of a gene chip, in virto as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence is that of a nucleic acid sequence of the invention. Note: The sequence data for this patent did not form part of the printed segmentally, but was obtained in electronic format directly from WIPO
                                                              This invention relates to novel isolated nucleic acid sequences involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; NAC-1; NAC-1 associated disorder; drug addiction; cancer; probe;
                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+03; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 5 A; 2 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                             ftp.wipo.int/pub/publishedpct_sequences
                                Disclosure; SEQ ID NO 261; 30pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NAC-1 gene-specific probe #150.
                                                                                                                                                                                                                                                                                                                                                                                                                                         2380 CATCTTGCCTCCAGGT 2395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 CATCTTGCCTCCAGAT
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                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
 transfected cells
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es 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ZHAN/) ZHANG J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1870
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Matches
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
IKappa Kinase (IKK), or protein Kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (ONS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), injury, spinal cord injury, cancer (e.g. melanoma,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased calls or to detect the presence of
target RNA in a cell. The present RNA sequence represents a human NOGO
                                                                                                                                                                                                      antisense oligonucleotide, neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PRR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimnume disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhintis; atopic dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fosnaugh K;
                                                                                                                                                                   Human NOGO receptor inozyme substrate sequence #324.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                receptor inozyme substrate sequence.
                                         ВЪ.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-APR-2002; 2002WO-US010512.
                                         ADL46891 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-058513/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200281628-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified.
                                                                                                                           20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-0CT-2002.
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                                                                                   ADL46891;
RESULT 1871
                      ADL46891
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The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a harmerhead ribozyme, NGH ribozyme, G-cleaver ribozyme, Zinzyme, Dinzyme, Dinzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, reducing GRID activity in a cell by contacting a patient having a condition associated with the level of GRID (e.g. tissue/graft rejection or leukaemia) by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by cortexing the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                               Human; 88; GRID; Grb2-related with insert domain; hammerhead ribozyme; NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme; hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.4%; Score 14.4; DB 1; Length 17; Best Local Similarity 75.0%; Pred. No. 1.6e+03; Matches 12; Conservative 3; Mismatches 1; Indels
                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hamblin PA,
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         Pred. No. 1.6e+03;
81.2%; Pred. ....
                                                                                                                                                                                                                                                                                               Human GRID mRNA substrate sequence #645.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 645; 74pp; English.
                                                                                                                                                                                       ADM54335 standard; mRNA; 17 BP.
                                                          1312 GATGCCACTGACAAGG 1327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-FEB-2001; 2001US-00792818.
                                                                                1 GAUGCCGCUGACAAGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-FEB-2000; 2000US-0181594P.
                                                                                                                                                                                                                                                           (first entry)
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                        13; Conservative
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CARLOWITZ I V.
MCSWIGGEN J.
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ELLIS J H.
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   Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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(ELLI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (JARV/)
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(MCSW/)
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The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a hammer-back inbozyme, NGT ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell ncluding the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID creating a patient having a condition associated with the level of GRID contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule in amanner that allows its expression, a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acide of the invention.
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                                                                                                                                                                                                                                                                       Human; ss; GRID; Grb2-related with insert domain; hammerhead ribozyme;
NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme;
hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
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                                                                                                                                                                                                                                    Human GRID mRNA substrate sequence #382.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 382; 74pp; English.
2006 TGGTGGAGGACCTGGA 2021
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                    1 UGGUGGAGGUCCUGGA 16
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                                                                                                                            ADMS4107 standard; mRNA; 17
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                                                                                                                                                                                                  (first entry)
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CARLOWITZ I V.
MCSWIGGEN J.
HAMBLIN P A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (JARV/) JARVIS
(CARL/) CARLOWI
                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                              ADM54107;
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(HAMB/)
                                                                                       RESULT 1873
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Gaps ö BP.

ADM59278/c ID ADM59278 standard; RNA; 17

ADM59278;

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New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                    Human, 88; GRID, Grb2-related with insert domain, hammerhead ribozyme; NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme; hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
                                                                                                                                                                                                                                                                                                               Ellis JH;
                                                                                                                                                                                                                                                                                                             Mcswiggen J, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 2 A; 2 C; 9 G; 0 T; 4 U; 0 Other;
                                                                                Human GRID mRNA substrate sequence #877.
                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 880; 74pp; English.
                    ADMS4567 standard; mRNA; 17 BP
                                                                                                                                                                                                        23-FEB-2001; 2001US-00792818.
                                                                                                                                                                                                                              10-FEB-2000; 2000US-0181594P.
                                                            (first entry)
                                                                                                                                                                                                                                                                                                              Jarvis T, Carlowitz IV,
                                                                                                                                                                                                                                                (JARV/) JARVIS T.
(CARL/) CARLOWITZ I V.
(MCSW/) MCSWIGGEN J.
(HAMB/) HAWBLIN P A.
(ELLI/) ELLIS J H.
                                                                                                                                                                                                                                                                                                                                  WPI; 2003-829646/77
                                                                                                                                                                 US2003134806-A1.
                                                                                                                                              Homo sapiens.
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                                                            03-JUN-2004
                                       ADM54567;
                                                                                                                                                                                                                                                                                                                                                                                      leukemia.
RESULT 1874
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The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNZxyme, amberzyme, Inozyme or hairpin ribozyme, Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID (e.g. tissue/graft rejection or leukaemia) by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at least the novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.
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0.4%; Score 14.4; DB 1; Length 17; 75.0%; Pred. No. 1.6e+03; ive 3; Mismatches 1; Indels
                                                        2006 TGGTGGAGGACCTGGA 2021
           75.0%;
                             12; Conservative
                Local Similarity
 Query Match
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Matches
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RESULT 1875

RESULT 1876 ADM57907/c

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The invention relates to an enzymatic nucleic acid molecule that specifically cleaves RNA derived from hepatitis B virus (HBV) and comprising one or more binding arms, without requiring the presence of a 2'-0H group within the molecule for activity. The nucleic acids are useful for treating hepatitis B virus infection, hepatitis, hepatocellular carcinoma, cirrhosis and liver failure, either alone or in combination with other therapies such as lamivudine and interferons. The nucleic acids are useful as diagnostic tools to examine genetic drift and mutations within diseased cells, for detecting the presence of HBV RNA in a cell, for the study of RNA and for down-regulating gene expression of target genes in bacterial, fungal, viral, plant or mammalian cells. This sequence represents an HBV RNA target sequence, used in the scope of the invention. Note: The sequence data for this patent is also available in electronic format from USPTO at sequata.uspto.gov/sequence.html.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel enzymatic nucleic acid molecule such as DNAzymes and inozymes specifically cleaving RNA derived from hepatitis B virus and comprising one or more binding arms, useful for treating hepatitis and cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                           Hepatitis B virus, HBV, ss, enzymatic nucleic acid; RNA cleavage, hepatitis B virus infection, hepatitis, hepatocellular carcinoma; cirrios; liver failure, lamivudine, interferon, genetic drift; virucide, hepatotropic, antinflammatory; cytostatic.
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                                                                         Hepatitis B virus (HBV) RNA target sequence #1412.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Draper K, Blatt L, Mcswiggen JA, Morrissey D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 1412; 122pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-2000; 2000US-00531025.
09-AUG-2000; 2000US-00635385.
24-OCT-2000; 2000US-00695347.
08-JUN-2001; 2001US-00877478.
                                                                                                                                                                                                                                                                                                                    15-JAN-2003; 2003US-00342902
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99US-00436430
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                                      03-JUN-2004 (first entry)
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MCSWIGGEN J A.
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Matches 15; Conserv
                                                                                                                                                                                                        Hepatitis B virus.
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                                                                                                                                                                                                                                            US2004054156-A1.
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08-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                        14-MAY-1992;
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Novel enzymatic nucleic acid molecule such as DNAzymes and inozymes specifically cleaving RNA derived from hepatitis B virus and comprising one or more binding arms, useful for treating hepatitis and cirrhosis.
                                                                                                                                   Hepatitis B virus; HBV; ss; enzymatic nucleic acid; RNA cleavage; hepatitis B virus infection; hepatitis; hepatocellular carcinoma; cirrhosis; liver failure; lamivudine; interferon; genetic drift; virucide; hepatotropic; antinflammatory; cytostatic.
                                                                                            Hepatitis B virus (HBV) RNA target sequence #2089.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 2089; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen JA,
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94US-00193627.
99US-00436430.
2000US-00531025.
2000US-00636385.
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                                                         03-JUN-2004 (first entry)
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Matches 10; Conservative
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MCSWIGGEN J A.
MORRISSEY D.
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                                                                                                                                                                                                                                      Hepatitis B virus
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07-FEB-1994;
08-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comparising one or more binding arms, without requiring the presence of a 2'-OH group within the molecule for activity. The nucleic acids are useful for treating hepatitis B virus infection, hepatitis, hepatitis, hepatitis, not the arctinoma, cirrhosis and liver failure, either alone or in combination with other therapies such as lamivudine and interferons. The nucleic acids are useful a diagnostic tools to examine genetic drift and mutations within diseased cells, for detecting the presence of HBV RNA in a cell, for the study of RNA and for down-regulating gene expression of target genes in bacterial, fungal, viral, plant or mammalian cells. This sequence represents an HBV RNA target sequence, used in the scope of the invention. Note: The sequence data for this patent is also available in electronic format from USPTO at sequence.oy/sequence.html.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel enzymatic nucleic acid molecule such as DNAzymes and inozymes specifically cleaving RNA derived from hepatitis B virus and comprising one or more binding arms, useful for treating hepatitis and cirrhosis.
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                                                                                                                                                      Hepatitis B virus; HBV; ss; enzymatic nucleic acid; RNA cleavage; hepatitis B virus infection; hepatitis; hepatocallular carcinoma; cirrhosis; liver failure; lamivudine; interferon; genetic drift; virucide; hepatotropic; antiinflammatory; cytostatic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.4%; Score 14.4; DB 1; Length 17; 33.8%; Pred. No. 1.6e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 3 A; 7 C; 5 G; 0 T; 2 U; 0 Other;
                                                                                                                 Hepatitis B virus (HBV) RNA target sequence #41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Morrissey D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 41; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                               14-MAY-1992; 92US-008B2712.
07-FEB-1994; 94US-00193627.
08-NOV-1999; 99US-00436430.
20-MAR-2000; 2000US-00531025.
09-AUG-2000; 2000US-0063885.
24-OCT-2000; 2000US-00636347.
08-JUN-2001; 2001US-00877478.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3626 GGCCCTGAGTCTGGG 3641
BP.
                                                                                                                                                                                                                                                                                                                                                                              15-JAN-2003; 2003US-00342902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93.88;
ADM57907 standard; RNA; 17
                                                                            (first entry)
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ses 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MCSW/) MCSWIGGEN J A. (MORR/) MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Draper K, Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-247781/23.
                                                                                                                                                                                                                                                         Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DRAPER K.
                                                                                                                                                                                                                                                                                                US2004054156-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BLATT L.
                                                                            03-JUN-2004
                                                                                                                                                                                                                                                                                                                                       18-MAR-2004.
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16
                                       ADM57907
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(BLAT/)
(MCSW/)
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Morrissey D;

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The invention relates to an enzymatic nucleic acid molecule that specifically cleaves RNA derived from hepatitis B virus (HBV) and comprising one or more binding arms, without requiring the presence of a 2'-OH group within the molecule for activity. The nucleic acids are useful for treating hepatitis B virus infection, hepatitis, hepatocellular carcinoma, cirrhosis and liver failure, either alone or in combination with other therapies such as lamivudine and interferons in nucleic acids are useful as diagnostic tools to examine genetic drift and mutations within diseased cells, for detecting the presence of HBV RNA in a cell, for the study of RNA and for down-regulating gene expression of target genes in bacterial, fungal, viral, plant or mammalian cells. This sequence represents an HBV RNA target sequence, used in the scope of the invention. Note: The sequence data for this patent is also available in electronic format from USPTO at sequata.uspto.gov/sequence.html.
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Pred. No. 1.6e+03;
5; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 4 A; 4 C; 3 G; 0 T; 6 U; 0 Other;
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ID AAX672
XX
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ADM59955 standard; RNA; 17 BP

RESULT 1877

ADM59955 ID ADM5

Matches

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16 ACAGGGGGCCAGGCATG 1

(first entry)

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The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising; (i) at least 5 ribose residues; (ii) a 2-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, oparticularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for chancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene cypression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is solved that their required of antieness molecules, and is highly the consentration are required of antieness molecules, and is highly
                                                                                                         Arthritic condition, graft tolerance, immune response; target; cleavage; hammerhead ribozyme; halrpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ensymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases.
                                                                        Mouse CD40 hairpin ribozyme target SEQ ID NO:3872.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 2 A; 9 C; 3 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 10; Page 219; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-300653/30.
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Mcswiggen J,
Karpeisky A,
                                                                                                                                                                                      diagnosis; ss
                                     20-JUL-1999
                                                                                                                                                                                                                                                                WO9618736-A2
                                                                                                                                                                                                                                                                                                                                      22-NOV-1995;
                                                                                                                                                                                                                                                                                                 20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                            13-DEC-1994;
23-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-1994;
17-FEB-1995;
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02-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-AUG-1995
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940S-00363254. 950S-00436850. 950S-00432874. 950S-00434509. 950S-0000951P.

95US-00512861 95US-00541365

95WO-US015516 94US-00354920 94US-00363253 Stinchcomb DT, Jarvis T, Draper K, Pavco P; Gustofson J, Usman N, Wincott F, Matulic-Adamic J; Thompson JD, Modak A, Burgin A;

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AAV02684-V02687 are PCR primers used to amplify the human leukocyte antigen (HLA) class I HLA-C gene intron 1 for locus-specific antigen (HLA) class I HLA-C gene intron 1 for locus-specific complication which is used in a novel method for testing a tissue sample to determine the allelic type of a HLA class I gene in the sample. The C HLA class I gene is selected from among HLA-A, -B and -C genes. The method comprises of treating the tissue sample to obtain nucleic acid comprises of treating the tissue sample to obtain nucleic acid comprises suited the complex suited a portion of intron 1 or intron 3 of the HLA class I gene and a second primer which hybridises with a confiterent portion of the HLA class I gene under conditions suitable for amplification to obtain an amplified product. The product is then evaluated to determine the allelic type of the HLA-class I gene. The certain suseful for tissue matching HLA class I antigens between donors and recipients and hence for preventing graft versus host disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - useful for
                                                                                                                                                      Human leukocyte antigen class I gene; HLA-C; allele testing; donor;
tissue matching; recipient; graft rejection; class typing; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA-based human leukocyte antigen class I gene typing method tissue matching and prevention of graft versus host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 18 BP; 2 A; 7 C; 9 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                     SLOK ) SLOAN KETTERING INST CANCER RES
                                                                                                                      Human HLA-C gene intron 1 PCR primer 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 13; Page 14; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1412 CGCAGGGCGGCCCCT 1427
                AAV02686 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV48450 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                  96WO-US000362.
                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cereb N;
                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                  04-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                    04-JAN-1996;
                                                                                                                                                                                                                                                                              WO9723645-A1
                                                                                     19-MAY-1998
                                                                                                                                                                                                                                                                                                                 03-JUL-1997.
                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yang SY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV48450;
                                                 AAV02686;
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AAV48450/C
ID AAV48450
XX
AC AAV48450
XC
XX
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AAV02686,
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Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels

15-0CT-1998 (first entry)

Gaps

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The present sequence is used in the exemplification of the

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This is the nucleotide sequence of CUT1 gene downstream primer cutprol. It is homologous to an upstream region of the Arabidopsis thaliana CUT1 gene (see AAV84182), and was used with upstream primer cutprol (see AAV84182) to amplify a 1949 bp fragment (see AAV84184) of the CUT1 gene promoter, and with upstream primer cutprol (see AAV84184) to amplify a truncated (1209 bp) promoter. These promoter fragments were used in promoter. The results indicate that the CUT1 promoter is regulated in a tissue-specific and cell-specific manner, and that epidermis specificity is retained in unrelated plant species such as tobacco. No differences in the strength of expression were detected between the 1.9 kb and 1.2 kb promoter. The CUT1 promoter can be used to produce transgene constructs that are specifically expressed in epidermal cells. The CUT1 gene encodes
                                                                                                                                                                                                                                                                                                                                                                                      encoding protein with very long chain fatty acid elongase activity - 1 to create transgenic plants with modified very long chain fatty acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   that are specifically expressed in epidermal cells. The ČUT1 gene encodes a protein (see AAW84180) involved in very long chain fatty acid synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zollinger-Ellison syndrome; gastrointestinal ulceration; congenital microvillus atrophy; skin disease; cell growth; abnormal keratinoryte differentiation; psoriasis; epithealial cancer; Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin; dermal scarring; Usher Syndrome; Atrophia areata; anti-thrombotic; wound healing; tissue repair; PCR primer; 86.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Secreted protein; transmembrane protein; human; enterocolitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; cive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Forward PCR primer used to amplify cDNA encoding PRO294.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 4 A; 2 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 11; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3106 GGCGGAGAGTTTTAAT 3121
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                                                                                                                                                                              97US-0043831P.
98US-00058947.
                                                                                                                                                                                                                                                (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                      98WO-CA000343.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15; Conservative
Arabidopsis thaliana.
                                                                                                                                                                                                                                                                                               Millar AA;
                                                                                                                                                                                                                                                                                                                                          WPI; 1999-080740/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 15; Conserv
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                                               WO9846766-A1
                                                                                                                                                                                   14-APR-1997;
                                                                                                                                      14-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 composition.
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                                                                                         22-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                               Kunst L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX52417;
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  셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transforming growth factor beta-1 (TGF beta-1). The oligonucleotides transforming growth factor beta-1 (TGF beta-1). The oligonucleotides exemplify the invention. The specification describes oligonucleotides that contain 8-30 mucleotides, which contain at most 8 mucleotides that consecutive mucleotides but to form three H-bonds each to four consecutive mucleotides able to form three H-bonds each to four consecutive cytosines, do not contain two sequences of three consecutive mucleotides able to form three H-bonds each to four three such bonds (3R) is given by 2R/3R = 0.33-0.72. The cytosines are used to modulate expression of genes, particularly the grammar of primary cell cultures (e.g. bone marrow stem, liver or kidney cells, osteoclasts, osteoblasts and/or keratinocytes). The coligonucleotides can also be used to analyse function of proteins (by coligonucleotides can also be used to analyse function of proteins (by caltering their expression or activity) and therapeutically, e.g. in cases of cancer or (targeting TGF) for stimulating the immune system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                          Transforming growth factor beta-1; TGF beta-1; antisense oligonucleotide; modulate; gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CUII; very long chain fatty acid elongase; VLCFA; wax; transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
  Transforming growth factor beta-1 antisense oligonucleotide N38.
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                                                                                                                                                                                                                                                                                                                                                                 (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CUT1 gene promoter downstream primer cutprol.
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                                                                                                                                                                                                                                                                                                                                                                                                              Schlingensiepen K, Brysch W;
                                                                                                                                                                                                                                                                          97EP-00101531
                                                                                                                                                                                                                                                                                                                       97EP-00101531
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nes 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-400910/35
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                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                          31-JAN-1997;
                                                                                                                                                                                                                                                                                                                       31-JAN-1997;
                                                                                                                                                                                   EP856579-A1
                                                                                                                                                                                                                               05-AUG-1998
                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV84183;
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16-SEP-1998;

17-SEP-1997; 17-SEP-1997; 17-SEP-1997; 17-SEP-1997; 17-SEP-1997;

17-SEP-1997; 17-SEP-1997; 18-SEP-1997; 18-SEP-1997; 17-OCT-1997; 17-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997; 24-OCT-1997;

27-0CT-1997; 28-0CT-1997; 28-0CT-1997; 28-0CT-1997; 28-0CT-1997; 28-0CT-1997; 29-0CT-1997; 29-0CT-1997;

29-OCT-1997; 29-OCT-1997; 29-OCT-1997; 29-OCT-1997; 31-OCT-1997;

31-OCT-1997; 03-NOV-1997; 07-NOV-1997;

17-NOV-1997 21-NOV-1997 21-NOV-1997

12-NOV-1997 18-NOV-1997 25-NOV-1997;

24-NOV-1997 24-NOV-1997

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diseases associated with abnormal keratinocyte differentiation (e.g. psoriaais, epithelial cancers such as lung squamous cell carcinoma of the vulva and gliomas), potent effects on cell growth and development, diseases related to growth or survival of nerve cells including Parkinson's disease, Alzheimer's disease, ALS, neuropathies or cancer. PRO26s can be used as for fibromodulin, e.g. for reducing dermal scarring. PRO264 can be used as a target for anti-tumor drugs. PRO533 may be used in the treatment of Usher Syndrome or Atrophia areata; PRO269 can be used as an anti-thromobic agent, PRO287 polypeptides and portrions may have therapeutic applications in wound healing and tissue repair; PRO317 can be used for treating problems of the kidney, uterus, endometrium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This oligonucleotide is complementary to a portion of the 5' untranslated region and start codon of the human G protein coupled receptor flh845 sea encorresponding to nucleotides 766-783 of the sequence given in AAX59167. It can be used to modulate flh845 activity, and hence to treat a disease or disorder characterized by, or associated with, aberrant or abnormal flh845 nucleic acid expression and/or flh845 polypeptide activity by inhibiting flh846 nucleic acid expression and/or flh845 polypeptide activity by inhibiting flh846 nucleic acid expression. Diseases and disorders associated with aberrant or abnormal flh845 activity include nervous system related disorders, e.g. ammestia, apraxia, agnosia, ammestic dysnomia, amnestic spatial disorientation, Kluver-Bucy syndrome,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid encoding an isolated G-protein coupled receptor useful
gastrointestinal ulceration and congenital microvillus atrophy), skin
                                                                                                                                                                                                                                                          blood vessels, or related tissue, e.g. in the heart of genital tract
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G protein coupled receptor; flh84g5; human; diagnosis; screening; therapy; antiparkinsonian; nootropic; neuroprotective; neuroleptic; antidepressant; antiarrythmic; antidiabetic; antiinflammatory; phosphatidylinositol; antisense; ss.
                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.4; DB 1; Length 18; 13.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human flh84g5 gene 5' region antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Goodearl ADJ, Glucksmann MA, Xie M, Distefano P;
                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         treating nervous system related disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 64; 140pp; English. /
                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                        1101 GCTGTCCTCAGGGGAG 1116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX59173 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98WO-US025832.
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98US-00042780.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 GCTGTCCACAGGGAG 18
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                                                                                                                                                                                                                                                                                                                                                                      93.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 93.8
les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-394858/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9928470-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX59173;
                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotides AAX52276-532 represent PCR primers and probes used to isolate and amplify cDNA encoding secreted and transmembrane human proteins (see AAX52213-74 and AAY1344-403). The CDNA sequences are obtained from cDNA libraries, prepared from fetal lung, fetal kidney, fetal brain, fetal liver and fetal retina. The encoded polypeptides have specific uses based on their homology to known polypeptides, e.g. PRO211 and PRO217 can be used for disorders associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions (e.g. enterocolitis, Zollinger-Ellison syndrome,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated human genes and polypeptides used in, e.g. treatment of gastrointestinal ulceration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 37; Page 143; 320pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9705-0063542P
9705-0063544P
9705-0063550P
9705-0063735P
9705-0063735P
9705-0063732P
9705-0063734P
9705-0063734P
9705-0063734P
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970S-0066466P.
970S-0066511P.
970S-0066770P.
970S-0066772P.
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97US-0062816P.
97US-0063045P.
97US-0063121P.
97US-0063127P.
97US-0063127P.
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97US-0059184P.
97US-0059263P.
97US-0059266P.
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97US-0062285P.
97US-0062287P.
97US-0063486P.
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97US-0063329P.
97US-0063541P.
                                                                                 97US-0059117P.
97US-0059119P.
97US-0059121P.
98WO-US019330
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wood WI, Gurney AL,
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Affancing consciousness such as visual hallucinations, purcelled affecting consciousness such as visual hallucinations, perceptual disturbances or delerium associated with Lewy body dementia, schizophrenia with mood swings, depressive illness conferences or schizophrenia with mood swings, depressive illness (primary and secondary); affective disorders such as REM sleep abnormalities in patients suffering from e.g. depression, paradoxical aleap abnormalities, sleep-wakefulness, and body temperature or respiratory depression abnormalities during sleep; disorders affecting pain generation mechanisms e.g. parkinson's disease related cor chest pain; movement disorders e.g. parkinson's disease related movement disorders, e.g. diabetic polydipsis, smooth muscle cobssity or drinking disorders, e.g. diabetic polydipsis, smooth muscle cobssity or drinking disorders, e.g. diabetic polydipsis, smooth muscle curinary incontinence, ossophageal achalasia or chronic obstructive arways disease; cardiac muscle disorders, e.g. pathologic bradycardia or tachycardia, arrhythmia, flutter or fibrillation; and gland related
                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat; muscarinic acetylcholine receptor 6; mACHR-6; detection; antidabetic; antidaressant; anciropic; neuroprotective; neuroleptic; antidiabetic; antidepressant; antiarrhythmic; antiinflammatory; carnitine; pain; G-protein coupled receptor; nervous system related disorder; xerostomia; disorders affecting consciousness; affective disorder; movement disorder; irritable bowel syndrome; drinking disorder; gland related disorder; smooth muscle related disorder; cardiac muscle disorder; eating disorder; diabetes mellitus; diagnosis; drug screening; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid encoding an isolated G-protein coupled receptor useful for treating nervous system related disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
  Alzheimer's related memory loss and learning disability; disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat mACHR-6 antisense oligonucleotide SEQ ID NO:24.
                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 2 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                          disorder such as xerostomia or diabetes mellitus
                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Col 49; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3658 GCCTGCAGGGCCATGG 3673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH44579 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GCCTGCTGGGCCATGG 16
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98US-00042780
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                                                                                                                                                                                                                                                                                                                                                                                                                     93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAR-2003 (revised)
01-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-394858/33.
                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Glucksmann MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH44579;
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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The present invention describes muscarinic acetylcholine receptor 6 (mACHR-6), which is a member of the G family of proteins. mACHR-6 has

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antidepressant, antiarrhythmic and antiinflammatory activities. The mACHR corp actide antidepressant, antiarrhythmic and antiinflammatory activities. The mACHR corp action, is capable of modulating the effects of a G-protein coupled receptor (GPCR) ligand such as activities or an acetylcholine like colorule such as carnitine, e.g. by modulating phospholipase C signalling/activity. Products from the present invention can be used for treating disorders mediated by abnormal mACHR of protein activity such as nervous system related disorders, disorders affecting consciousness, affective disorders such as REM sleep abnormalities, disorders affecting consciousness, affective disorders such as pain related to irritable bowell syndrome or chest pain, movement disorders, eating disorders, drinking disorders, smooth muscle related disorders, cardiac muscle disorders, and gland related disorders such as xerostomia or diabetes mellitus. The present asquence represents a rat mACHR-6 antisense oligonucleotide which is given in the exemplification of the present invention. (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method (I) for testing a tissue sample to determine the allelic type of a human leukocyte antigen (HLA) class I gene in the sample, where the HLA class I gene is selected from HLA-A, HLA-B or HLA-C, by: (a) treating the tissue sample to obtain nucleic acid polymers suitable for amplification; (b) combining the nucleic acid polymers with a primer which hybridizes with a portion of intron 1 or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Testing a tissue sample to determine the allelic type of a human leukocyte antigen class I gene comprises amplification of nucleic acid polymers with primers which flank a region including an allelic variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue sample testing, allelic typing; human leukocyte antigen; PCR primer; probe; hybridisation; intron; amplification; 88; allelic variation; non-classical HLA class I gene; exon.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Locus specific amplification primer #3 for HLA-C gene.
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                  on 20-MAR-2003 to correct DR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 13; Col 11; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3658 GCCTGCAGGCCATGG 3673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA11070 standard; DNA; 18 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GCCTGCTGGGCCATGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                      93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUL-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polymers with primers wh
of the HLA class I gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-223159/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CERE/) CEREB N. (YANG/) YANG S Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-DEC-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA11070;
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intron 3 of the HLA class I gene, and a second primer which hybridizes with a different portion of the HLA class I gene and performing amplification, where the primers flank a region including at least one site of allelic variation in at least one of exons 2 or 3 of the HLA class I gene and where the first primer is a locus specific primer which hybridizes with intron 1 or 3 of only one of the HLA class I genes; and (c) evaluating the amplified product to determine the allelic type of the HLA class I gene. The method is useful for testing a tissue sample to determine the allelic type of a classical or non-classical HLA class I gene in the sample. The sequences AAAI1039-AAI112 represent consensus sequences of introns and exons of the HLA genes and primers and probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiinflamatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; vasotropic; chemotaxic; angiogenic; neuroprotective; vasotropic; chemotaxic; antiarthritic; antirheumatic; antiarthritic; antirheumatic; antiarthritic; antirheumatic; antiarthritic; antirheumatic; thrombolyric; immunomodulator; enterocolitis; collinger-Ellison syndrome; gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease; Alzheimer's, ALS; neuropathy; dermal scarring; wound healing; esthma; theumatoid arthritis; multiple sclerosis; inflammatory disorder; atherosclerosis; cardiac injury; infertility; premeture aging; AlDS; diabetes; stroke; gene therapy; transgenic; PRO; human; ss; primer; PCR.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PRO protein-related forward PCR primer SEQ ID 229.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 2 A; 7 C; 9 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                  used to isolate and analyse the HLA genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1412 CGCAGGGCGGCCCCT 1427
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Matches
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scarring and wound healing, nerve repair, thrombosis, bone and/or cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple sclerosis, inflammatory disorders, atherosclerosis, cardiac injury, infertility, premature aging, AIDS, diabetes complications and stroke. The molecules may also be utilised during gene therapy procedures and transgenic animal production. The current sequence is that of the PCR primer of the invention which was used to analyse the human PRO DNA of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sixty one nucleic acids encoding PRO polypeptides which are useful in the
                                                                                                                                                                                                                                                                                                                                                                                                               Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory; antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant; antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer; antiarthritic; antiinfertility; antidiabetic; antiviral; diabetes; ophthalmological; gene therapy; skin disease; gastrointestinal disorder; ischaemia; inflammation; PCR primer; ss.
                                                                                                                                                                                  Gaps
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ME, Goddard A
Kljavin IJ;
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0
                                                                                                                                                     Score 14.4; DB 1; Length 18;
Pred. No. 1.7e+03;
                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide gene PCR primer SEQ ID NO: 229.
                                                                                                                            Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                          0.4%; Scc...
v 93.8%; Pred. No. 1...
... 0; Mismatches
                                                                                                                                                                                                            1101 GCTGTCCTCAGGGGAG 1116
                                                                                                                                                                                                                                                                                                        AAF72575 standard; DNA; 18 BP
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99US-0146222P.
99WO-US020594.
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99WO-US030095.
99WO-US030911.
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                                                                                                                                                                                  15; Conservative
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                                                                                                                                                       Query Match
Best Local Similarity
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Filvaroff E,
                                                                                                  the invention.
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05-OCT-1999;
29-NOV-1999;
30-NOV-1999;
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                                                                                                                                                                                                                                                                               RESULT 1887
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Novel nucleic acids encoding secreted and transmembrane polypeptides with homology, e.g. to growth and cancer-associated antigens.

Example 37; SEQ ID NO 229; 355pp; English.

Wood

Pennica D,

Gurney AL, Hillan K,

98WO-US019330

16-SEP-1998;

(GETH) GENENTECH INC. Goddard A,

WPI; 2000-271434/23.

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Chen Yuan The invention relates to a novel nucleic acid encoding a PRO polypeptide. The polypeptides and polynucleotides of the invention may be useful as research tools and as therapeutics for treating enterocolitis. Sollinger-Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer, Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal

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29-OCT-1997;
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                                                                                                           invention
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                                                                                                                                                         Query Match
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24-0CT-1
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27-0CT-1
27-0CT-1
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28-0CT-1
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ACA60188
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                                                                   The present sequence is a primer which was used in the isolation of one of sixty one nucleic acids encoding novel secreted and transmembrane PRO polypeptides. The PRO polypeptides are useful for treating skin diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma), gastrointestinal disorders (e.g. enterocolitis), neurodegenerative diseases (e.g. Alzheimer's disease, Parkinson's diseases, wound repair, cardiovascular disorders (e.g. endometrial bleeding angiogenesis, ischaemias such as coronary ischaemia, atherosclerosis), inflammatory disorders (e.g. asthma, rheumatoid archritis, multiple sclerosis), inflammatory infertility, AIDS and diabetes and retinal disorders such as retinitis pigmentcoum. The PRO nucleic acids have applications in molecular biology, including use as hybridization probes, and in chromosome and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a collection of binding groups for a family of nucleic acids comprising members of relative high and relative low significance, where the binding groups are selected to be capable to identify, alone or in combination, essentially all members of the family of nucleic acids of relatively high significance. The collection of
ment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Collection of binding groups for determining or typing samples, especially clinical samples, has groups capable to identify essentially all members of the family of nucleic acids of relatively high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 related binding molecule oligonucleotide sequence SEQ ID NO:24.
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                                                                                                                                                                                                                                                                 Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            molecule; HIV-1; human immunodeficiency virus type transcriptase; binding group; ss.
                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                        Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                0.4%; Score 14.4; DB 1; 93.8%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                        0; Mismatches
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                                               Example 37; Page 181; 393pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                 GCTGTCCTCAGGGGAG 1116
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                                                                                                                                                                                                                                                                                                                                                                                                 ABL88802 standard; DNA; 18
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 treatment of skin
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                                                                                                                                                                                                                    gene mapping
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                       disease)
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                     sample,
binding groups is useful for typing of nucleic acid in a clinical sample by contacting the nucleic acid with the collection and determining whether one or more binding groups bound to the nucleic acid of the sample. This method is useful for determining whether the sample comprises at least a part of a member of relatively high significance or a family of nucleic acids. The collection of binding groups is useful fidagnosing the severity of a disease caused by a pathogen containing a member of a family of nucleic acids. ABL88779 to ABL89321 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03;
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97US-0059115P.
97US-0059117P.
97US-0059121P.
97US-0059122P.
97US-0059184P.
97US-0059263P.
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97US-0063435P.
97US-0063704P.
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Matches 15; Conservative
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vivlemore401-10.rng

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16-SEP-1998; 98W0-US019330.
17-SEP-1998; 98W0-US019437.
01-DEC-1999; 99W0-US025108.
08-SEP-1999; 99W0-US020944.
13-SEP-1999; 99W0-US020944.
15-SEP-1999; 99W0-US021647.
05-OCT-1999; 99W0-US023089.
05-OCT-1999; 99W0-US023089.
01-DEC-1999; 99W0-US028564.
02-DEC-1999; 99W0-US028565.
16-DEC-1999; 99W0-US028565.
16-DEC-1999; 99W0-US028565.
16-DEC-1999; 99W0-US028565.
16-DEC-1999; 99W0-US028565.
16-DEC-1999; 99W0-US03099.
05-JAN-Z000; 2000W0-US003565.
22-FEB-Z000; 2000W0-US003565.
22-FEB-Z000; 2000W0-US003565.
22-FEB-Z000; 2000W0-US003565.
22-FEB-Z000; 2000W0-US0056641.
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30-MAR-2000; 2000WO-US008439.
22-MAY-2000; 2000WO-US014042.
02-JUN-2000; 2000WO-US015264.
28-JUL-2000; 2000WO-US02710.
24-AUG-2000; 2000WO-US023328.
18-SEP-2000; 2000US-00665350.
                    97US-0063734P
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(GETH) GENENTECH INC.

Goddard A: Kljavin IJ; Tumas D; Ferrara N; Gerber H, Gerritsen Mrney AL, Hillan KJ, Roy MA, Stewart TA, Eaton DL, Ashkenazi A, Botstein D, Desnoyers L, Pilvarcoff B, Pong S, Gao W, Gerber H, Godowski PJ, Grimaldi JC, Gurney AL, Mather JP, Pan J, Paoni NF, Roy MA, an J, Pa Wood WI; Mather JP, P

WPI; 2003-329602/31.

New transmembrane polypeptides and nucleic acids encoding the polypeptides, useful in gene therapy, in chromosome identification, as chromosome markers, in generating probes and in tissue typing.

Example 37; Page 111; 484pp; English.

nucleic acid sequence identity to a nucleotide sequence encoding one of 61 secreted/transmembrane polypeptides, or PRO polypeptides or encoding a PRO protein extracellular domain. Also included are a vector comprising the PRO mucleic acid, a host cell comprising the vector, producing a PRO polypeptide (by culturing the host cell for the expression of the PRO polypeptide, and recovering the PRO polypeptide from the cell culture), The invention relates to an isolated nucleic acid with at least 80%

17-OCT-1997; 17-OCT-1997;

18-SEP-1997 15-OCT-1997

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an isolated PRO polypeptide (having at least 80% sequence identity to: (
a) an amino acid sequence selected from the 61 PRO proteins; (b) an amino
acid sequence encoded by a nucleic acid molecule deposited with an ATCC
number (detailed in the specification); or (c) an extracellular domain of
a PRO polypeptide or to a PRO polypeptide lacking its associated signal
CC peptide), a chimaeric molecule comprising a PRO polypeptide of fused to a
cherrologous amino acid sequence, an anti-PRO antibody, detecting a
cp PRO145 or PRO1668 in a sample suspected of containing the polypeptide,
cc protection and sequence, an anti-PRO antibody, detecting a
cmodulating at least one biological activity of a cell expressing a PRO245
cor PRO1868. Nucleic acids which encode PRO can be used to generate either
cc transgenic animals or knock-out animals which may be used in the
development and screening of therapeutically useful reagents. The nucleic
acids may also be used in gene therapy, in chromosome identification, as
chromosome markers, or in generating probes. The PRO polypeptides are
cc useful as molecular markers for protein electrophoreals, and the isolated
cnucleic acids may be used for recombinantly expressing those markers. The
course of polypeptides and nucleic acids may also be used in diagnostic assays for PRO, and in
calfinity purification of PRO from recombinant cell culture or natural
csurces. The present sequence is a PCR primer used to isolate a cDNA
control of PRO protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; secreted and transmembrane protein; PRO; pharmaceutical; diagnostic, biosensor; bioreactor; Parkinson's disease; allemmation; nephritis; wound healing; nerve repair; collateral blood vessel formation; cancer; colorectal cancer; haemorrhage; rheumatoid arthritis; diabetes; cirrhosis; fibrosis; restenosis; dermal fibrotic condition; keloid; scarring; ischaemia; stroke; hypertension; heart attack; atherosclerosis; infertility; gene therapy; PCR; primer; ss.
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02-DEC-1999
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Gurney AL, Botstein D, Des Fong S, Gao W, Grimaldi JC, Gu Ashkenazi A, Filvaroff E, Godowski PJ,

Desnoyers L, Eaton DL, Ferrara N; W, Gerber H, Gerritsen ME, Goddard A; Gurney AL, Hillan KJ, Kljavin IJ;

The invention describes a new isolated minimal and with the American Type Culture Collection (e.g. ATCC Deposit No. 209258), or a sequence of the DNA encoding a PRO polypeptide comprising any of 61 sequences having 164-1119 amino acids fully defined in the Specification. The PRO polypeptides or polymucleotides are useful as particularly useful for detecting or treating e.g. Parkinson's disease, particularly useful for detecting or treating e.g. Parkinson's disease, Alabeimer's disease, inflammations, nephritis, wound healing, nerve repair, collateral blood vessel formation, cancers (e.g. colorectal cancer), haemorrhage (or reduce risk for haemorrhage), rheumatoid arthritis, diabetes, cirrhosis of the liver, fibrosis of the lungs, carbritis, diabetes, cirrhosis of the liver, fibrosis of the lungs, cirrhosis, dermal fibrotic conditions (e.g. keloids or scarring), cirrhosis, dermals (e.g. humans, odgs, cats, cattle, horses, sheep, pigs, goats, or rabbits) The PRO polypeptides are useful as targets for theorems. The presence of these diseases. The PRO polypeptides are also useful as molecular weight markers, or for chromosome identification. The PRO genes are useful as hybridisation probes, or for screening libraries of the numan conna, probes, or for screening libraries of the numan conna, probability and carbridge may also be used in gene therapy, particularly for replacing a defective gene. This sequence created and transmembrane PRO polypeptide ö New genes and secreted and transmembrane polypeptides (e.g. PRO245 or PRO315), useful for treating or diagnosing e.g. Alzheimer's disease, cancers, hemorrhage, rheumatoid arthritis, diabetes, cirrhosis, ischemia The invention describes a new isolated nucleic acid molecule comprising Gaps .. 0 Stewart TA, Tumas D; Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Pan J, Paoni NF, Roy MA, wood WI; Example 37; Page 103; 482pp; English. 1101 GCTGTCCTCAGGGGAG 1116 WPI; 2003-370793/35 Mather JP, P. Williams PM, or strokes. 8

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BP.

ABX71636 standard; DNA; 18

RESULT 1891

10-MAR-2003 (first entry)

ABX71636;

Human, PRO; secreted protein, transmembrane protein, enterocolitis, gastrointestinal ulceration; skin disease; ss; PCR; primer; abnormal keratinocyte differentiation, psoriasis; epithelial cancer; sequamous cell carcinoma, Alzheimer's disease; Parkinson's disease; amyotrophic lateral sclerosis; inflammatory disease; rheumatoid arthritis; asthma; multiple sclerosis; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; disbetic complication; wound repair. Human secreted/transmembrane protein PRO294 PCR primer #2.

Ното варіелв.

US2002132240-A1

19-SEP-2002

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9703-0063735P.
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10-SEP-1998
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New PRO polypeptides and nucleic acid molecules, useful in diagnosing or treating inflammatory diseases, organ failure, atherosclerosis, cardiac injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's
                                                                                              Ferrara N;
ME, Goddard A;
Kljavin IJ;
                                                                                                                      Tumas D;
                                                                                            Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Williams PM, Wood WI;
      2000MO-US005841.
2000MO-US007377
2000MO-US014042.
2000MO-US01584.
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                                                                              (GETH ) GENENTECH INC.
                                                                                                                                             WPI; 2003-147434/14.
      02-MAR-2000; 20-MAR-2000; 230-MAR-2000; 22-MAY-2000; 20-JUN-2000; 28-JUL-2000; 2
                                                              18-SEP-2000;
                                                      24-AUG-2000;
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Example 37; Page 101; 473pp; English

The invention relates to an isolated PRO polypeptide having at least 80% amino acid sequence identity to: (a) any one of 61 fully defined amino acid sequences given in the specification (appearing as AB19347-CAP164407); (b) an amino acid sequence encoded by the nuclectide sequence acid sequence encoded by the nuclectide sequence acid sequence encoded by the nuclectide sequence listed in the specification); (c) any one of the PRO sequences which care is associated signal peptide; (d) any one of the PRO sequences which compared to stracellular domain of the PRO polypeptide with its associated signal peptide; (d) an extracellular domain of the PRO polypeptide with its associated signal peptide. Also include are the nucleic acids encoding the PRO polypeptides vectors, host cells and anti-PRO antibodies. The PRO polypeptides and nucleic acids are useful in diagnosing or treating enterocolitis, gastrointestinal ulceration, skin diseases associated with abnormal kerstaincyte differentiation, e.g. psoriasis or epithelial cancers such as squamous cell carcinoma, Alaheimer's disease, e.g. cherosclerosis, cardiac injury, infertility, birth defects, premature arthritis, asthma or multiple sclerosis, or goneral. The concerned with re-growth of tissue. The mucleotide sequences may be used concerned with re-growth of tissue. The mucleotide sequences may be used to an tybridisation probes in chromosome and gene mapping, or in generating antipolypeptides, in assays to identify other proteins or molecules involved in binding reaction, to generate transgenic animals or knockout animals, which in turn are useful in the development and screening of therapeutically useful reagents, for chromosome identification, and clerrophoreais purposes. The anti-DRO mile markers for protein concerned and seasys for or for the affinity purification or protein concerned and sample or an area of the process or protein or protein

recombinant cell culture or natural sources. The present sequence is a PCR primer used to isolate a cDNA encoding a PRO polypeptide Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ; 0 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels 15; Conservative Best Local Similarity Matches 15; Conserv Query Match

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1101 GCTGTCCTCAGGGGAG 1116 18 GCTGTCCACAGGGGAG m

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hysterectomy; angiogenesis; coronary ischaemic condition; skin disease; gastrointestinal mucosa disorder; acute mucosal lesion; neuropathy; ALS; chronic mucosal lesion; abnormal keratinocyte differentiation; psoriasis; Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis; parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis; harcointeid edil growth, cancer; blood cosqulation cascade; thrombosis; haemorrhage; endometrial bleeding; angiogenesis; wound healing; asthma; tissue repair; rheumatoid arthritis; multiple sclerosis; tissue typing;
                                                                   Human; PCR; primer; abnormal bleeding; gynaecological disease; tumour;
                                                     Human secreted/transmembrane polypeptide PRO294 forward primer #2
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                                      (first entry)
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29-OCT-1997;
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29-OCT-1997;
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29-0CT-1997;
31-0CT-1997;
31-0CT-1997;
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24-NOV-1997;
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                       ACH06968;
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97US-0066770P.
97US-0066470P.
97US-0066840P.
98US-0099802F.
98UG-0099803F.
98UG-0100262F.
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99US-0143048P.
99US-0145698P.
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98US-0100858P.
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99WO-US021090
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25-NOV-1997;
12-DEC-1997;
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15-SEP-1
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(GETH) GENENTECH INC.

Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvarcif E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; Williams PM, Wood WI;

WPI; 2003-492258/46.

Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating abnormal bleeding involved in gynecological diseases, skin diseases and neurodegenerative diseases.

Example 37; Page 107; 478pp; English.

The invention relates to an isolated PRO polypeptide. PRO317 is useful in diagnosing or treating abnormal bleeding involved in gynecological diseases e.g. to avoid or lessen the need for hysterectomy. PRO317 may also be useful as an agent that affects angiogenesis and PRO317 is useful in anti-tumour indications or in treating coronary ischeamic conditions. PRO311 and PRO217 polypeptides are useful for treating disorders associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte differentiation (e.g. psoriasis). PRO387 polypeptide is useful for treating Parkinson's disease, Alzheimer's disease, amyotrophic lateral sclerosis (ALS), neuropathies

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New secreted and transmembrane PRO polypeptides (e.g. PRO533 or PRO245)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Botstein D, Desnoyers L, Eaton DL, Ferrara N;
Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
an J, Paoni NF, Roy MA, Stewart TA, Tumas D;
                970S-0063128P
970S-006332P
970S-0063541P
970S-0063542P
970S-0063542P
970S-0063564P
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99WO-US021090.
99WO-US021547.
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24-AUG-2000; 2000WO-US023328.
18-SEP-2000; 2000US-00665350.
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98WO-US019437.
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2000WO-US005841
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11-FEB-2000;
22-FEB-2000;
24-FEB-2000;
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Filvaroff E,
Godowski PJ,
     24-0CT-1997;
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20-MAR-2000;
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05-OCT-19
29-NOV-19
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31-0CT-19
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       and disease related to uncontrolled cell growth, e.g. cancer. PR0219

CC polypeptide plays a regulatory role in the blood coagulation cascade.

PR0246 polypeptides which serves as tumour specific antigens may be exploited as therapeutic targets for anti-tumour drugs. PR0269 exploited as therapeutic targets for anti-tumour drugs. PR0266 polypeptide is useful as an antithrombotic agent with reduced risk for haemorrhage as compared with heparin. PR0317 polypeptide is useful in creating endomerial bleeding angiogenesis. PR0287 polypeptides and polypeptides and polypeptides and polypeptides and tissue repair. PR0214 polypeptides are useful for treating asthma, rheumatoid arthritis, psoriasis and multiple sclerosis. The polypeptide and tissue repair. CC are useful for tissue typing. PRO antibodies are useful for cassay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its cypuspide are useful in diagnostic assays for PRO e.g. detecting its cypuspide cells, tissues or serum and for affinity present sequence represents a human secreted/transmembrane PRO cyc polypeptide PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; PCR; primer; ss; PRO; secreted; transmembrane; pharmaceutical; diagnostic; biosensor; bioreactor; therapeutic; hyperplassia; endometriosis; cancer; tumour; ischaemia; coronary arterial disease; polycytic kidney disease; temal failure; inflammatory response; asthma; rheumatoid arthritis; psoriasis; multiple sclerosis; gene therapy; cytostatic; gynecological; cardiant; nephrotropic; hepatotropic; antiinflammatory.
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                                                                                                                                                                                                                                                 0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted/transmembrane protein, #42, PCR primer #2.
                                                                                                                                                                                                                            Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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97US-0059113P.
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97US-0063814P.
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Best Local Similarity 93.8
Matches 15; Conservative
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21-0CT-1997;
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24-FEB-2000;
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05-OCT-1999;
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17-NOV-1997
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        The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. The PRO polypeptides or polymucleotides are also useful as pharmaceuticals, diagnostics, biosensors or bioreactors, for detecting or treating e.g. hyperplasia, endometriosis, cancers (e.g. those involving solid tumours), ischaemia, coronary arterial disease, polyyostic kidney disease, chronic or acute renal failure, or inflammatory responses (e.g. asthma, rheumatoid arthritis, psoriasis or multiple sclerosis) in mammals. The PRO genes may gene. The sequences presented in ABX96017-ABX96378 are the genes encoding, the primers amplifying and the probes detecting the PRO polynucleotides of the invention
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and genes encoding them, useful for detecting or treating e.g.
hyperplasia, endometriosis, cancers, ischemia, coronary arterial disease
or inflammations.
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                                                              Example 36; Page 107; 477pp; English
                                                                                                                                                                                                                                                                                                                                                                                                    1101 GCTGTCCTCAGGGGAG 1116
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97US-0062287P.
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les 15; Conservative
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98WO-US019177.
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03-NOV-1
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 The invention relates to sixty one nucleic acids encoding PRO

CD polypeptides (secreted and transmembrane). The polynucleotide is useful

CC in molecular biology, including uses as hybridisation probes, in

CC chromosome and gene mapping, in generating antisense RNA and DNA, and in

CC gene therspy. The polynucleotide may also be used in preparing PRO

CC gene therspy. The polynucleotide may also be used in preparing PRO

CC polypeptides by recombinant techniques, and in generating either

CC transgenic animals or knock-out animals which, in turn, are useful in the

CC polypeptide or the antibody is used in preparing a medicament for

CC polypeptide or the antibody is used in preparing a medicament for

CC polypeptide or the antibody is used in preparing a medicament for

CC treating a condition responsive to the polypeptide or antibody, such as

mucosal lesions e.g. ulcers and enterocolitis, skin disease e.g.

C psoriasis, cancer e.g. lung cancer and colon cancer, nerve cell disease

CC atrophia areata, adisease and Parkinson's disease, Usher syndrome,

CT atrophia areata, andiogenesis, inflammatory disease e.g asthma and

C rheumatoid arthritis, ischaemia, and in various diagnostic assays. The

CC polypeptide

CC polypeptide
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                                                                                                                                                                                                                                     e.g. PRO245
                                                                                                                                                                                                                                    Sixty one isolated nucleic acids encoding a PRO polypeptide, e.g. PRO2 or PRO1868, useful in chromosome and gene mapping, in generating antisense RNA and DNA, and in treating cancer and Alzheimer's disease.
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                                                                                       Ferrara N;
ME, Goddard A;
Kljavin IJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted / transmembrane polypeptide PRO294 forward primer.f2.
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                                                                                                                                                Tumas D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; Ive 0; Mismatches 1; Indels
                                                                                       Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NF, Roy MA, Stewart TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                              Example 37; Page 110; 481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1101 GCTGTCCTCAGGGGAG 1116
24-AUG-2000; 2000WO-US023328.
18-SEP-2000; 2000US-00665350.
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Best Local Similarity 93.8%;
Matches 15; Conservative
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                                                                                     Ashkenazi A, Botstein E
Filvarcff B, Fong S, G
Godowski PJ, Grimaldi J
Mather JP, Pan J, Paon
Williams PM, Wood WI,
                                                   (GETH ) GENENTECH INC.
                                                                                                                                                                                                   WPI; 2003-331485/31.
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02-DEC-1999; 99WO-US028564.

02-DEC-1999; 99WO-US028565.

20-DEC-1999; 99WO-US030095.

20-DEC-1999; 99WO-US03011.

20-DEC-1999; 99WO-US03011.

20-DEC-1999; 99WO-US03011.

11-FEB-2000; 2000WO-US001565.

22-FEB-2000; 2000WO-US001565.

24-FEB-2000; 2000WO-US005841.

20-MAR-2000; 2000WO-US00841.

30-MAR-2000; 2000WO-US008439.

22-MAY-2000; 2000WO-US0084397.
                                                                            2000WO-US015264
2000WO-US020710
                                                                                         2000WO-US023328
                                                                                              2000US-00665350
                                                                                                          (GETH ) GENENTECH INC.
                                                                                                                                 Godowski PJ, Grin
Mather JP, Pan J,
                                                                                                                     Ashkenazi A,
Filvaroff E,
                                                                             02-JUN-2000;
                                                                                   28-JUL-2000;
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Ferrara N; n ME, Goddard A; Kljavin IJ; Tumas D; Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NP, Roy MA, Stewart TA, Wood WI: Williams PM,

WPI; 2003-417923/39

Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers.

Example 37; Page 106; 469pp; English.

The invention relates to an isolated, secreted and transmembrane polypeptide, termed PRO polypeptide. The polypeptide, for preparing variants of the polypeptide, so metagonists of the polypeptide is useful for preparing variants of the polypeptide. The polypeptide is also useful for preparing variants of the polypeptide is also useful as therapeutic electrophoreals purpose and the nucleic acid is useful for recombinantly expensing those markers. The polypeptide is also useful as therapeutic agent. PRO is useful in assays to identify other proteins or molecules involved in binding interaction. The nucleic acid is useful as therapeutic probes, in chromosome and gene mapping, in generation of antistense RNA and DNA, in the preparation of PRO pypeptide, for generating transgenic animals or knockout animals which in turn are construct hybridisation probes for mapping, in generation construct hybridisation probes for mapping with genetic asceders, in gene therapy, for chromosome identification, as chromosome marker, and for generating probes for polymerase chain reaction (PCR), overthern analysis of individuals which encodes the PRO and for the genetic analysis. PRO antibody is useful in diagnostic assays for PRO, e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from the preparation of medicament for treating conditions which is responsive to the PRO polypeptide or anti-PRO antibody eig tumour. The polypeptide and the nucleic acid is useful for the preparation of medicament typing. The polypeptide is useful for treating bone and/or cartilage or inhibiting tumour growth, enhances vascular permeability and immune treating hearing loss in mammals and for treating bone and/or cartilage creating hearing loss in mammals and entrating properior creating properior and arthiting the present engances. disorders such as sports injuries and arthritis. The present sequence represents a human secreted and transmembrane PRO polypeptide PCR primer

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

ö Gaps ö 0.4%; Score 14.4; DB 1; Length 18; 13.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels Query Match 0.4%; Best Local Similarity 93.8%; 15; Conservative Matches

3 GCTGTCCACAGGGGAG 18

셤

ABX11857 standard; DNA; 18 RESULT 1896

ABX11857;

Human muscarinic acetylcholine receptor 6 antisense oligonucleotide 10-MAY-2003 (first entry)

delirium; schizo-effective disorder; schizophrenia; depression; affective disorder; sleep disorders; pain generation disorder; irritable bowel syndrome; chest pain; movement disorder; Parkinson's disease; eating disorder; insulin hypersecretion obesity; heart muscle disorder; bradycardia; tachycardia; arrhythmia; flutter; fibrillation; gland related disorder; xerostomia; diabetes mellitus. cognitive disorder; amnesia; amnestic spatial disorientation; Kluver-Bucy syndrome; Alzheimer's related memory loss; antisense; learning disability; consciousness disorder; visual hallucination; Human; 88; mACHR-6; muscarinic acetylcholine receptor-6;

Ношо варіепв.

JS2002166131-A1.

07-NOV-2002

08-JUL-1999;

97US-00985090. 98US-00042780. 04-DEC-1997; 17-MAR-1998;

(MILL-) MILLENNIUM PHARM INC.

Glucksmann MA; WPI; 2003-298709/29. Goodearl ADJ,

New muscarinic acetylcholine receptor 6 (mACHR-6) nucleic acids and proteins, useful for modulating acetylcholine or phosphatidylinositol, particularly for treating e.g. schizophrenia, chest pain, tachycardia or arrhythmia.

Disclosure; Page 26; 66pp; English.

The invention relates to an isolated human or rat muscarinic acetylcholine receptor 6 (mACHR-6) nucleic acid molecule and the encoded protein. Also included are (non-human) host cells comprising the mACHR-6 nucleic acid molecule, an antibody that selectively bind the polypeptide above, a method for producing the polypeptide by culturing the host cell above, a method for buckler acid is expressed, a method for detecting the presence of the mACHR-6 polypeptide and nucleic acid, a method for detecting a compound that binds to the mACHR-6 polypeptide and a method for identifying a compound that binds to the mACHR-6 polypeptide and a method for identifying a compound that binds to the mACHR-6 polypeptide and a method for identifying a compound that binds to the mACHR-6 polypeptide. The mACHR-6 polypeptide and a method of compounding assays, diagnostic assays for identifying diseases, allelic planmacogenetic testing, methods of treatment, pharmacogenetic testing, methods of treatment, pharmacogenetic testing, methods of treatment, pharmacogenetic testing, polypeptide or antibody is useful carticular, the mACHR-6 polymuclectide, polypeptide or antibody is useful for treating or diagnosing cognitive disorders (e.g. ammessic, amnessic or treatment alsorientation, Kluver-Bucy syndrome, Alzheimer's related memory loss or learning disability), disorders (e.g. pain related to irritable bowel syndrome, or chest pain), movement disorders (e.g. enlisophers) and related disorders (e.g. parkinson's disease), eating disorders (e.g. insulin hypersecretion obesity), heart muscle related disorders (e.g. bradycardia, tarthythmia, flutter or fibrillation), or gland related disorder (e.g. arrhythmia, flutter or fibrillation), or gland related disorder (e.g. arrhythmia, elutter or fibrillation), or gland related disorder (e.g. arrhythmia, flutter or fibrillation), or gland related disorder (e.g. arrhythmia, flutter or fibrillation), plant related disorder (e.g. arrhythmia, arrhythmia, flutter or fibrillation).

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New PRO polypeptides and nucleic acid molecules, useful in diagnosing or treating inflammatory diseases, organ failure, atherosclerosis, cardiac injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's disease.
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02-MAR-2000;
20-MAR-2000;
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22-FEB-2000;
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Williams PM,
                       31-OCT-1997;
03-NOV-1997;
07-NOV-1997;
12-NOV-1997;
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18-NOV-1997
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ö
                                                                                                                                                                                                  Human; secreted and transmembrane protein; gene therapy; psoriasis; enterocolitis; gastrointestinal ulceration; skin disease; keratinocyte differentiation; epithelial cancer; Alzheimer's disease; squamous cell carcinoma; Parkinson's disease; inflammatory disease; amyotrophic lateral sclerosis; rheumatoid arthritis; asthma; multiple sclerosis; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; premature aging; AIDS; cancer; diabetic complication; wound repair; tissue re-growth; PCR; primer; ss.
                                                    Gaps
                                                                                                                                                                                   Novel secreted and transmembrane protein associated primer #103.
                                                    ö
                                0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; tive 0; Mismatches 1; Indels
                 Sequence 18 BP; 2 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
oligonucleotide targetting human mACHR-6
                                                                    3658 GCCTGCAGGGCCATGG 3673
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97US - 0062287P.
97US - 0063486P.
97US - 0063814P.
97US - 0062814P.
97US - 0063112P.
97US - 0063121P.
97US - 0063121P.
97US - 0063127P.
97US - 0063127P.
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97US-0059117P.
97US-0059117P.
97US-0059121P.
97US-0059122P.
97US-0059184P.
97US-0059263P.
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97US-0063544P.
97US-0063549P.
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97US-0063564P.
97US-0063435P.
97US-0063704P.
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97US-0063734P.
97US-0063735P.
                                                                               GCCTGCTGGGCCATGG 16
                                                                                                                                                                                                                                                                                                                                   2001US-00903640
                        Query Match
Best Local Similarity 93.00,
Conservative
                                                                                                                               ACA54996 standard; DNA; 18
                                                                                                                                                                  05-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                                 US2003017463-A1
                                                                                                                                                                                                                                                                                 Homo sapiens,
                                                                                                                                                                                                                                                                                                                                   11-JUL-2001;
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21-0CT-1997;
24-0CT-1997;
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Ferrara N;
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Kljavin IJ;
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97US-0064215P.
97US-0064248P.
97US-0064103P.
97US-0065180F.
97US-00651846P.
97US-0065634P.
97US-0065634P.
97US-0066421P.
97US-0066451P.
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2000WO-US004414.
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99WO-US021547
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18-SEP-2000; 2000US-00665350
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2000WO-US020710
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ashkenazi A, Botste
Filvaroff E, Fong S
Godowski PJ, Grimal
Mather JP, Pan J,
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The invention describes sixty one nucleic acids encoding PRO polypeptides (secreted and transmembrane). The PRO polypeptides and nucleic acids are useful in diagnosing or treating enterocolities, gastrointestinal ulceration, skin disease associated with abnormal keratinocyte differentiation, e.g. psoriasis or epithelial cancers such as squamous cell carcinoma, Alzheimer's disease, Parkinson's disease, amyotrophic lateral sclerosis, inflammatory diseases, e.g. rheumatoid arthritis, asthma or multiple sclerosis, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, cancer, canburt, infertility, birth defects, premature aging, AIDS, cancer, disbetic complications, or mutations in general. The polypeptides are also useful for wound repair and associated therapies concerned with respect to useful in gene therapy, and as molecular weight markers for protein electrophoresis purposes. The anti-PRO antibodies may be used in diagnostic assays for PRO, or for the affinity purification of PRO from tecombinant cell culture or natural sources. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphic site; nootropic; neuroprotective; anticonvulsant; anorectic; hypotensive; cardiant; thrombolytic; antiarteriosclerotic; osteopathic; antirheumatic; antiarthritic; antianflammatory; antinnfertility; psychiatric disorder; bipolar affective disorder; unipolar depression; gchizophrenia; anxiety; neurological disorder; obesity; insomnia; addiction; neurodegeneration; hypotension; hypertension; acute heart failure; atherothembosis; atherosclerosis; osteoporosis; rheumatoid arthritis; infertility; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polynucleotides encoding GPRSO receptor proteins and having at least one polymorphic site, useful for screening for GPRSO modulators for treating psychiatric disorders, e.g. bipolar affective disorder or unipolar depression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          novel human PRO polypeptide associated primer
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                                    Example 37; Page 101; 473pp; English.
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The present invention describes a polynucleotide sequence (I) which encodes a G protein-coupled receptor designated GPR50 and has at least encodes a G protein-coupled receptor designated GPR50 and has at least come polymorphic site. Also described are GPR50 allait variant polynucleotide sequences (ABZ81152 to ABZ81158) which encode the proteins given in ABR39074 to ABR39080. (I) has nootropic, neuroprotective, anticonvulsant, anorectic, properties, antiarthritic, antiarthritic, antiarteriosclerotic, osteopathic, antirheumatic, antiarthritic, antiarteriors and expression vectors from the present invention can be used in screening assays for identifying new drugs, and screening for USR50 modulators for preparing a medicament for treating psychiatric GRR50 modulators for preparing a medicament for treating psychiatric disorders, e.g. bipolar affective disorder or unipolar depression. They care also useful for correcting, preventing or ameliocating depression, schizophrenia, anxiety, neurological disorder, obseity, insomnia, addiction, neurodegeneration, hypotension, hypertension, acute heart atherothromboasis, atherosclerosis, osteoporosis, rheumatoid arthritis and infertility. The present sequence represents a PCR primer used to amplify the single nucleotide polymorphism (SNP) 1804 of human GPR50, which is used in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; 88; PCR; primer; gene therapy; apoptosis; bleeding; tumour; ALS; gynacclogical disease; hysterectomy; angiogenesis; skin disease; cancer; coronary ischaemic condition; gastrointestinal mucosa disorder; asthma; mucosal lesion repair; keratinocyte differentiation; psoriasis; Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis; neuropathy; blood coagulation cascade disorder; thrombosis; hemorrhage; neurodegenerative disease; endometrial bleeding; wound healing; tissue repair; rheumatoid arthritis; multiple sclerosis; tissue typing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 7 A; 5 C; 1 G; 5 T; 0 U; 0 Other;
                  Example 7; Page 20; 84pp; English.
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97US-0059115P.
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Best Local Similarity 93.8
Matches 15; Conservative
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97US-0062814P.
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26-0CT-1997;
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17-SEP-1998;
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05-OCT-1999;
29-NOV-1999;
30-NOV-1999;
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25-NOV-1997;
12-DEC-1997;
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13-OCT-1998;
20-NOV-1998;
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24-NOV-1997;
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26-JUL-1999;
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08-SEP-1999;
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17-NOV-1997
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The invention relates to an isolated secreted and transmembrane PRO CC polypeptide. The PRO polypeptides are useful for modulating biological activity of a cell, in diagnoseing or treating abnormal bleeding involved in gynaecological diseases e.g. to avoid or lessen the need for hysterectomy, for treating angiogenesis, tumour, coronary ischaemic condition, disorders associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte (differentiation) (e.g. psoriasis), Parkinson's disease, Alzheimer's disease, amyotrophic lateral sclerosis (ALS), neuropathies, disease condensing disorders, neurodegenerative disease, thrombosis, haemorrhage, endometrial bleeding, wound healing, tissue repair, asthma, rheumatoid arthritis, multiple sclerosis. Nucleic acid encoding PRO polypeptides are useful in molecular blology including uses as hybridisation probes and in the generation of antisense RNA and DNA, for preparing PRO polypeptides, conservation of antisense RNA and DNA for preparing PRO polypeptides, conservation of antisense RNA and DNA for preparing PRO polypeptides, conservating transgenic animals or knockout animals. The PRO antibodies are useful for tissue typing. PRO antibodies are useful for tissue so serum and conservating the series and their machenical staining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for conservation of affinity purification of PRO for recombinant cell culture or natural consemendation becomes an human secreted and conservations.
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                                                                                                                                                                                                                                                                                                                                                             Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating abnormal bleeding involved in gynecological diseases, skin diseases and neurodegenerative diseases.
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                                                                                                                                                                                           Ferrara N;
ME, Goddard A;
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Kljavin IJ;
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                                                                                                                                                                    Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen P Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, F Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Williams PM, Wood WI;
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                 22-MAY-2000; 2000WO-US014042.
02-UUN-2000; 2000WO-US015264.
28-UUI-2000; 2000WO-US020710.
24-AUG-2000; 2000WO-US023328.
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les 15, Conservative
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cell growth; cancer; tumour; viral infection; neurodegenerative disease; antithrombotic agent; haemorrhage; endometrial bleeding angiogenesis; diffiney tissue; apoptoals; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nootropic; neuroprotective; cytostatic; virucide; anticoagulant.
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97US-0059119P.
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17-SEP-1998;
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28-0CT-1997;
29-0CT-1997;
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31-0CT-1997;
31-0CT-1997;
31-0CT-1997;
31-0CT-1997;
03-NOV-1997;
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98US-0104080P 98US-0103304P 98US-0113296P 99US-0143048P 99US-0145698P 99US-0145622P 99WO-US021547 99WO-US023089 99WO-US028214. 99WO-US028313. 99WO-US028301. 99WO-US020944. 2000WO-US000219. 99WO-US028564 99WO-US028565 99WO-US030095 99WO-US030911 99WO-US030999 2000WO-US003565 2000WO-US005841 2000WO-US008439 2000WO-US023328 2000US-00665350 2000WO-US004414 2000WO-US005044 24-FEB-2000; 02-MAR-2000; 15-SEP-1999; 05-OCT-1999; 29-NOV-1999; 05-JAN-2000; 11-FEB-2000; 22-FEB-2000; 02-JUN-2000; 20-DEC-1999;

(GETH) GENENTECH INC.

Ferrara N; ME, Goddard A; мы, Goddar Kljavin IJ; Eaton DL, Gerritsen Hillan KJ, Stewart TA, Botstein D, Desnoyers L, Fong S, Gao W, Gerber H, ldi JC, Gurney AL, Paoni NF, Roy MA, Fong S, Gao Grimaldi JC, Pan J, Pao 4, Wood WI; Ashkenazi A, Filvaroff E, Godowski PJ, Mather JP, P Williams PM,

WPI; 2003-765473/72.

Novel isolated native PRO polypeptide useful for treating Parkinson's disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal ulceration, Alzheimer's disease, amyotrophic lateral sclerosis, Usher syndrome

Example 36; Page 98; 469pp; English

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for linking a concertive molecule to a cell expressing a PRO proteptides are useful for detecting other PRO polypeptides in a sample and for linking a bloactive molecule to a cell expressing a PRO polypeptides are useful compared to a cell expressing a PRO polypeptide and activity of a cell expressing PRO polypeptides. PRO polypeptide are also useful compared to a cell expressing PRO polypeptides. PRO polypeptides are also useful conformation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte of differentiation (e.g. psoriasis), Parkinson's disease. Alzheimer's differentiation (e.g. psoriasis), Parkinson's disease. Alzheimer's differentiation (e.g. psoriasis), parkinson's diseases. Alzheimer's cadditionally, disease related to uncontrolled cell growth, e.g. cancer. PRO polypeptides also serves as tumour specific antigens which may be exploited as therapeutic targets for anti-tumour drugs, and are also erves as tumour specific antigens which may be exploited as therapeutically in vivo for lessening the effects of viral infection. The PRO polypeptides can be also used in assays to determine contitith and a roll in neurodegenerative diseases or their reversal, as an antithrombotic agent with reduced risk for haemorrhage as compared with heparin, in treating other PRO-associated disorders, in modulating endometrial bleeding angiogenesis, and may also have an effect on kidney

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97US-0063486P
97US-0062814P
97US-0063101P
97US-00631212P
97US-00631212P
97US-00631212P
97US-0063124P
97US-0063544P
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24-0CT-1997;
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24-0CT-1997;
27-0CT-1997;
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        tissue. PRO polypeptides and their portions affect the expression of genes which have a role in apoptosis. The polynucleotides are useful in and-cular biology including uses as hybridisation probes for cDNA library to isolate the full-length PRO cDNA or to isolate other cDNAs, in the generating transgenic animals or chromosome and gene mapping, in the generating transgenic animals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals which are useful in the development and screening of therapeutic agents, for as probes and for the genetic analysis of individuals with genetic disorders as well as for the proteins are useful are protein and for chromosome identification. The proteins are useful are useful as molecular marker for protein electrophoresis purposes, as charapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (agonists) or prevent the effect of the PRO cloppeptide (antagonists). The polynucleotides and proteins are useful collypeptide (and of or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific calls, itssues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The equence presented is a PCR primer which was used to amplify a PRO or polymention.
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93.8%;

Best Local Similarity

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20-MAR-2000;
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Matches RESULT 19 ACD66978 ò 셤 cand the nuclear discinges isolated by the polypeptides can be used to raise and the nuclear discinges isolated by the polypeptides can be used to raise inthodies that specifically bind to the PRO polypeptide. For linking a consideration biological activity of a call. PRO polypeptide. The PRO bioactive molecule to a call expressing a PRO polypeptide. The PRO bioactive molecule to a call expressing a PRO polypeptide. The PRO bioactive molecule to a call expressing a PRO polypeptide. The PRO polypeptide artibodies are useful for modulating the biological activity of a call expressing PRO polypeptides. PRO polypeptides are also useful for the consideration and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal cof gastrointestinal mucosa and the repair of acute and chronic mucosal cof gastrointestinal mucosa and the repair of acute and chronic mucosal cof gastrointestinal mucosa and the repair of acute and chronic mucosal cof gastrointestinal mucosa and the repair of acute and chronic mucosal cof gastrointestinal mucosa and the repair of acute and chronic mucosal cofficential properties a sociated with abnormal keratinocyte diseases anyotrophic lateral sclossis (ALS), neuropathies and calculation (e.g. psoriasis) Parkinson's disease, anyotrophic lateral sclossis (ALS), neuropathies and calculation. The PRO polypeptides can be also used in assays to determine infection. The PRO polypeptides can be also used in assays to determine if thas a role in neurodegenerative disorders in modulating cantity properties and their portions affect the expression of antitatory to isolate the full-length PRO cassociated disorders, in modulating of concernation probes for connecting and poptodis. The polymcloticides are useful in the generation of antisense RNA and DNA, to isolate the full-length PRO cassociated disorders, in modulating of the reparting and poptodis, or isolate the full-length PRO cassociated concernation of antisense RNA and DNA, to isolate the full-length PRO cassociated concernatio The invention discloses isolated PRO secreted/transmembrane polypeptides New secreted and transmembrane polypeptides useful for treating skin, neurodegenerative diseases, asthma, rheumatoid arthritis, psoriasis and multiple sclerosis. recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO ME, Goddard A; Kljavin IJ; Ferrara N; Tumas D; Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen erimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NF, Roy MA, Stewart TA, Example 36; SEQ ID NO 229; 471pp; English. polynucleotide of the invention. 30-MAR-2000; 2000WO-US008439. 22-MAY-2000; 2000WO-US014042. 02-JUN-2000; 2000WO-US015264. 28-JUL-2000; 2000WO-US020710. 24-AUG-2000; 2000WO-US023328. 18-SEP-2000; 2000US-00665350. Filvaron Gram Godowski PJ, Gram Fabra Fabra Wood WI; (GETH) GENENTECH INC. WPI; 2003-503392/47. Ashkenazi A, Filvaroff E,

0.4%; Score 14.4; DB 1; Length 18;

Query Match

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                                                                                                                                                        Human, 88; PRO; secreted and transmembrane protein; inflammation; rheumatoid arthritis; psoriasis; multiple sclerosis; atherosclerosis; infertility; birth defect, premature aging; malignancy; cancer; stroke; heart attack; hypertension; gastrointestinal ulceration; parkinson's disease; Alzheimer's disease; AIDS; cholesterol uptake; wound healing; tissue repair; gene therapy.
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                                                                                                                                        Human secreted/transmembrane protein PRO294 PCR primer #2.
        Indels
Pred. No. 1.7e+03;
0; Mismatches 1
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9705-0063486P
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97US-0064809P.
97US-0065186P.
                                                                                      ACD66978 standard; DNA; 18 BP
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97US-0062285P.
97US-0062287P.
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                                          GCTGTCCACAGGGGAG 18
                                                                                                                       (first entry)
        15; Conservative
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97US-0066120P.
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97US-0069425P.
98US-0088026P.
98US-0099803P.
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11-FEB-2000; 2000WO-US003565.
22-FEB-2000; 2000WO-US004414.
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20-MAR-2000; 2000WO-US007377.
30-MAR-2000; 2000WO-US008439.
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02-JUN-2000; 2000WO-US015264.
28-JUL-2000; 2000WO-US020710.
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18-SEP-2000; 2000US-00665350
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      18-NOV-1997;
21-NOV-1997;
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25-NOV-1997;
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10-SEP-1998;
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01-DEC-1999
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(GETH) GENENTECH INC.

ME, Goddard A; Kljavin IJ; Tumas D; Ferrara N; Desnoyers L, Eaton DL, W, Gerber H, Gerritsen Gurney AL, Hillan KJ, NF, Roy MA, Stewart TA, Ashkenazi A, Botstein D, Desnoyers L, Filvaroff E, Fong S, Gao W, Gerber H Godowski PJ, Grimaldi JC, Gurney AL, Mather JP, Pan J, Paoni NF, Roy MA, Williams PM, Wood WI;

WPI; 2003-512316/48.

New genes and secreted and transmembrane polypeptides (e.g. PRO245 or PRO1868), useful for treating or diagnosing e.g. cancers, atherosclerosis, infertility, stroke, AIDS or multiple sclerosis in mammals.

Example 37; Page 103; 476pp; English.

The invention relates to an isolated nucleic acid molecule comprising a agequence with at least 80% identity to: (a) a nucleotide encoding any of 61 PRO (secreted and transmembrane protein) polypeptides appearing as AB032756-AB032816; or (b) any of 61 nucleotide sequences having 50-4053bp

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Tury uncleased in the particularity, or the full rengular county bequences of any these of nucleotide sequences. Also included are the isolated PRO polypeptide (lacking its associated signal peptide or an extracellular domain of the PRO polypeptide, with or lacking its associated signal peptide or an extracellular comprising the vector comprising the nucleic acid molecule, a host cell comprising the vector (used to produce the PRO polypeptide or an extracellular comprising the PRO polypeptide fused to a heterologous amino acid sequence, an anti-PRO antibody, detecting PRO245 or PRO1868 polypeptides, inking a bioactive molecule to a cell expressing a PRO245 or PRO1868 polypeptide and modulating at least one biological activity of a cell expressing the PRO245 or PRO1868 polypeptide. The PRO polypeptides or polymucleotides are useful as pharmaceuticals, diagnostics, biosensors or polymucleotides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These are particularly useful for diagnostics, biosensors or bioreactors, rheese are particularly useful for diagnostics, malignary (e.g. cancers), strokes, heart attacks, hypertension, gastroincestinal ulcarations, Parkinson's diseases, Alzheimer's disease, or AlDS in the body, and in wound healing or tissue repair. The PRO polypeptides are useful as hybridisation probes, or for screening libraries of human consecutions in the propertic care useful as hybridisation probes, or for screening libraries of human consequence is an oligomucleotide (PCR primer or probe) used in the cestorion of a PRO CDNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              abnormal bleeding; gynaecological disease; hysterectomy; mucosal lesion; coronary ischaemic condition; gastrointeetinal mucosa; skin disease; ALS; keratinocyte differentiation; psoriasis; Parkinson's disease; asthma; Alzhaimer's disease; arheumatoid arthritis; multiple sclerosis; cancer; amyotrophic lateral sclerosis; neuropathy; uncontrolled cell growth; PCR.
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  or the full length coding sequence
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fully defined in the specification;
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ses 15; Conservative
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18-SEP-1997;
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97US-0063486P.
97US-0062814P.
97US-0062816P.
97US-0063045P.
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98US-0088026P.
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17-0CT-1997;
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29-0CT-1997;
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31-0CT-1997;
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29-NOV-1999;
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28-OCT-1997;
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17-SEP-1998;
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15-SEP-1999;
15-SEP-1999;
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The invention relates to human PRO polypeptides (secreted and transmembrane polypeptides) and the PRO polynucleotides encoding them. The PRO polypeptides and polynucleotides can be used in diagnosing or treating abnormal bleeding involved in gynaecological diseases e.g. to avoid or lessen the need for hysterectomy. They can also be used in treating coronary ischaemic conditions, disorders associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte differentiation (e.g. psoriasis), Parkinson's disease, Alzheimer's disease, asthma, rheumatoid arthritis, multiple sollerosis, amyotrophic lateral scherosis (ALS), neuropathies and diseases related to uncontrolled cell growth, such as cancer. This sequence represents a PCR primer used to isolate a human PRO polynucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neconatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; etimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritis; anorectic.
                                                                                                                                                                                                                                                                                                                                                                                                    Novel secreted and transmembrane PRO polypeptides and polynucleotides encoding them, useful for treating abnormal bleeding involved in gynecological diseases, skin diseases and neurodegenerative diseases.
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, Gerritsen ME, Goddard A;
Hillan KJ, Kljavin IJ;
Stewart TA, Tumas D;
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Fong S, Gao W, Gerber H,
Grimaldi JC, Gurney AL, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 37; Page 103; 475pp; English.
24-FEB-2000; 2000WO-US005004.
02-WAR-2000; 2000WO-US005841.
20-WAR-2000; 2000WO-US001437.
30-WAR-2000; 2000WO-US014042.
22-MAY-2000; 2000WO-US015264.
28-UJU-2000; 2000WO-US015264.
24-UJU-2000; 2000WO-US023328.
18-SEP-2000; 2000WS-0065350.
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Matches 15; Conservative
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Filvaroff B, Fong S, G
Godowski PJ, Grimaldi J
Mather JP, Pan J, Paon
Williams PM, Wood WI,
                                                                                                                                                                                                      (GETH ) GENENTECH INC.
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Homo sapiens

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99WO-US020944. 99WO-US023089 99WO-US028214, 99WO-US028313, 99WO-US028301, 99WO-US028564, 2000WO-US003565. 2000WO-US015264. 99WO-US021547 99WO-US028565 99WO-US030095 2000WO-US008439 2000WO-US014042 99WO-US030911 99WO-US030999 2000WO-US000219 2000WO-US005841 2000US-00665350 15-SEP-1999; 15-SEP-1999; 05-OCT-1999; 29-NOV-1999; 30-NOV-1999; 02-MAR-2000; 20-MAR-2000; 30-MAR-2000; 22-MAY-2000; 05-JAN-2000; 16-DEC-1999; 20-DEC-1999; 20-DEC-1999

(GETH) GENENTECH INC.

Ferrara N; ME, Goddard A; Kljavin IJ; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, E Filvaroff B, Fong S, Gao W, Gerber H, Gerritsen N Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, F Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Williams PM, Wood WI; WPI; 2003-521801/49.

New genes encoding for secreted and transmembrane PRO polypeptides, useful for treating e.g. cardiac insufficiency disorders, wounds, cancers, obesity, diabetes, hyperinsulinemia, hypoinsulinemia, or arthritis.

Example 36; SEQ ID NO 229; 476pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antithodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a call expressing a PRO polypeptide, for Inking a contection of the properties are useful contective molecule to a call expressing a PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a contective molecule to a call expressing a PRO polypeptide. The PRO polypeptides are useful for modulating the biological activity of a call expressing a properties. The PRO polypeptides or contective molecule to a call expressing a properties. The PRO polypeptides or polymented are useful as pharmaceuticals, diagnostics, biosensors or polymelectides are useful as paramaceuticals, diagnostics, biosensors or proliferation of endothelial calls growth factor (VEGF) stimulated proliferation of conforceceptor calls, inducing proliferation of conforceceptor calls, inducing proliferation of conforceceptor calls, inducing proliferation and/or recalls, modulating glucose or PRA uptake, inducing proliferation and/or recalls, modulating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to creinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a cole in cell death. The polymucleotides are useful in molecular biology including uses as hybridisation probabes or correspondent and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides and their portions and sor and some mapping, in the generation of antisense Contection animals which are useful in the development and contection and provides and for the polyment and sor kenting and provides and provides and provided the proportion and gene mapping, in the gene

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     recombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (agonists). The polynucleotides and proteins are immunohistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful for tassue typing, PRO antibodies are useful for antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
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keratinocyte differentiation; psoriasis; Parkinson's disease;
Alzheimer's diseases, amyotrophic lateral sclerosis; ALS; neuropathy;
cell growth; cancer; tumour; viral infection; neurodegenerative disease;
antithrombotic agent; haemorrhage; endometrial bleeding angiogenesis;
kidney tissue; apoptosis; therapeutic; tissue typing; kidney tissue; ammunohistochemical staining; gene therapy; nootropic; neuroprotective;
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Mather JP, Pan J,
                                                        Williams PM,
 Ashkenazi A,
Filvaroff E,
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WPI; 2003-755054/71.

Novel PRO polypeptides useful for treating Parkinson's disease, Alzheimer's disease, enterocolitis, Zollinger-Ellison syndrome, psoriasis, epidermoid carcinoma of the vulva and gliomas, gynecological diseases

Example 36; SEQ ID NO 229; 479pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides can define auclate acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO polypeptide, for inking at least one biological activity of a cell. PRO polypeptide, are useful for modulating at least one biological activity of a cell. PRO polypeptide are useful for determine the properties. The properties are activated to cell expressing a PRO polypeptide are useful for modulating the biological activity of a cell expressing a PRO polypeptide. The PRO polypeptide articological activity of a cell expressing and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual maintenance of gastrointesinal mucosa and the rapair of actual may and a daditionally, disease related to uncontrolled cell growth, e.g. cancer. PRO polypeptides also serves as tumour specific antigens which may be expension of maintenance of maintenance of maintenance and additionally, disease related to uncontrolled cell growth, e.g. cancer. PRO polypeptides and their portions affect the expression of material may also have an effect on kidney to isolate the full-leading angiogenesis, and may also have an effect on kidney to isolate the full. Leading angiogenesis, and may also have an effect on solate the full-leading angiogenesis, and may also have a role in apoptosis. The polymucleotides are useful in chargemine and properties and properties and their properties and propertie

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

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0.4%; Score 14.4; DB 1; Length 18; 13.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels
                       93.8%;
                     Best Local Similarity 93.8
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Human; PRO; primer, ss; Parkinson's disease; Alzheimer's disease; ALS; amyotrophic lateral sclerosis; neuropathy; cancer; viral infection; AIDS; Usher's syndrome; haemorrhage; enterocolitis; Zollinger-Ellison syndrome;
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97US-0064103P.
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                                                                                                  (first entry)
                                                                                                                                          Human PRO PCR primer #96
                                                                                                                                                                                                                                                                                                                            diabetic complication.
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Local Similarity
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17-0CT-1997;
21-0CT-1997;
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                                                         invention
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Matches
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I ME, Goddard A;
Kljavin IJ;
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rney AL, Hillan KJ,
Roy MA, Stewart TA,
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22-FBB-2000; 2000WO-US004414.
24-FBB-2000; 2000WO-US005004.
20-MAR-2000; 2000WO-US005841.
20-MAR-2000; 2000WO-US005841.
30-MAR-2000; 2000WO-US00841.
22-MAY-2000; 2000WO-US008439.
97US-0066511P.
97US-0066770P.
97US-0066840P.
97US-0069425P.
98US-0089026P.
98US-0099803P.
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22-DEC-1998;
07-JUL-1999;
26-JUL-1999;
28-JUL-1999;
24-NOV-1997;
24-NOV-1997;
25-NOV-1997;
12-DEC-1997;
10-SEP-1998;
10-SEP-1998;
14-SEP-1998;
14-SEP-1998;
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13-SEP-1999;
15-SEP-1999;
15-SEP-1999;
05-OCT-1999;
30-NOV-1999;
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17-SEP-1998;
17-SEP-1998;
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20-NOV-1998
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aging, AIDS, stroke and diabetic complications. The polynucleotides are also useful in chromosome and gene mapping. This sequence represents a PCR primer used in isolation of a human PRO polynucleotide of the
                                                                                                                     Gaps
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                                                                                          0.4%; Score 14.4; DB 1; Length 18;
llarity 93.8%; Pred. No. 1.7e+03;
Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                 Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                1101 GCTGTCCTCAGGGGAG 1116
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97US-0059113P
97US-0059121P
97US-0059121P
97US-0059122P
97US-0059128P
97US-006228SP
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97US-0063124P
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29-OCT-1997;
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97US-0063732P.
97US-0063734P.
97US-0063735P.
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11-FEB-2000; 2000WO-US003565
22-FEB-2000; 2000WO-US00414.
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02-JUN-2000; 2000WO-US015264
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18-NOV-1997;
21-NOV-1997;
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24-NOV-1997;
25-NOV-1997;
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(GETH) GENENTECH INC.

Ferrara N; 1 ME, Goddard A; Kljavin IJ; Tumas D; , Desnoyers L, bu. no W, Gerber H, Gerritu. rney AL, Hillan KJ, K. Ashkenazi A, Botstein D, Desnoyers L, Filvaroff B, Fong S, Gao W, Gerber F Godowski PJ, Grimaldi JC, Gurney AL, Mather JP, Pan J, Paoni NF, Roy MA, Williams PM, Wood WI,

WPI; 2003-503391/47.

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The invention discloses isolated PRO secreted/transmembrane polypeptides can durince acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptides are useful biological activity of a cell expressing a PRO polypeptides are useful concernive molecule to a cell expressing a PRO polypeptides are useful concernive molecule to a cell expressing a PRO polypeptides are useful concernive molecule to a cell expressing a PRO polypeptide and the PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a pRO polypeptide and maintenance of gastroinfestinal mucosa and the repair of acted informic mucosal for treating disorders associated with the preservation and maintenance of gastroinfestinal mucosa and the repair of acted informic mucosal lesions, skin diseases associated with abnormal keratinocyte differentiation (e.g. psoriasis), Parkinson's disease, Alizheimer's diseases, amyorrophic lesteral action for properties also serves as tumour specific antigens which may be disposed the arapeutic largets for anti-tumour drugs, and are also employed therapeutically in vivo for lessening the effects of viral conformation. The PRO polypeptides and under professor of microin and properties are useful in the arapid of activities for the part in the arapid properties and their professor of antithrombotic agent with reduced risk for hemorrhage as compared with have a role in apoptosis. The polymuclecides are useful in the development and screening of therapeutically useful reagents, as probes and for the generation of chromosome and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides, and their professor and proteins are useful in the development and screening of therapeutically useful reagents, as probes and for the generation of antisent and professor of individuals with genetic disorde
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secreted and transmembrane PRO polypeptides e.g. PRO187, which is per of the epidermal growth factor-8 (EGF-8) family of proteins,
                                                                                       Example 36; SEQ ID NO 229; 471pp; English.
                                           useful for treating cancer.
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Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

ö 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; rive 0; Mismatches 1; Indels Local Similarity 93.8 nes 15; Conservative Query Match Best Loca Matches

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Gaps

1101 GCTGTCCTCAGGGGAG 1116 3 gererecacadedad

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ADA13118 standard; DNA; 18 BP. 06-NOV-2003 (first entry) ADA13118; RESULT 1908 ADA13118 XEXEXEX

Human secreted/transmembrane protein, #44, PCR primer #2.

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gastrointestinal mucosa; mucosal lesion; skin disease; keratinocyte differentiation; psoriasis; Parkinson's disease; Alzheimer's diseases; amyotrophic lateral sclerosis; ALS; neuropathy; cell growth; cancer; tumour; viral infection; neurodegenerative disease; antithrombotic agent; haemorrhage; endometrial bleeding angiogenesis; kidney tissue; apoptosis; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nootropic; neuroprotective;
Human; PCR; primer; 88; PRO; secreted; transmembrane;
                                                                             cytostatic; virucide; anticoagulant.
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970S-0063121P
970S-0063122P
970S-0063329P
970S-0063542P
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970S-0063542P
970S-0063542P
970S-0063544P
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970S-0063564P
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97US - 0059113P .
97US - 0059113P .
97US - 0059121P .
97US - 0059124P .
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97US-0062287P.
97US-0063486P.
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97US-0062816P.
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98US-0100262P
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                                                                                                Homo sapiens.
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14-SEP-1998;
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Ferrara N;
1 ME, Goddard A;
Kljavin IJ;
Tumas D;
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                                      98WO-US025108.
98US-0113296P.
99US-0143048P.
99US-0145698P.
99US-0146222P.
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99WO-US028214.
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New secreted and transmembrane PRO polypeptides, useful for treating cancer, skin disorders, neurodegenerative diseases, and for lessening the effects of viral infection. WPI; 2003-521802/49.

Example 36; SEQ ID NO 229; 473pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for nuclating a conference of the properties and for linking a conference of conference of properties in a sample and for linking a conference of conference of properties are useful for discretive molecule to a cell expressing a PRO polypeptides. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptide are useful for modulating the biological activity of a cell expressing with the preservation and maintenance of conference of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte differentiation (e.g. psoriaais), parkinson's disease, Alzheimer's diseases, amyotrophic lateral sclerosis (ALS), neuropathies and additionally, disease related to uncontrolled cell growth, e.g. cancer. PRO polypeptides also serves as tumour specific antigens which may be exploited as therapeutic targets for anti-tumour drugs, and are also exploited as therapeutic targets for anti-tumour drugs, and are also comployed therapeuticapeutically in vivo for lessening the effects of viral ciffection. The PRO polypeptides can be also used in assays to determine if it has a role in neurodegenerative diseases or their reversal, as an

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            cc antithrombotic agent with reduced risk for haemorrhage as compared with heparin, in treating other PRO-associated disorders, in modulating conferencial bleeding angiogeneeis, and may also have an effect on kidney tissue. PRO polypeptides and their portions affect the expression of consensual biology including uses as hybridisation probes for cDNA library to isolate the full-length PRO cDNA or to isolate other cDNAs, in chromosome and gene mapping, in the generating transgenic animals or chromosome and gene mapping, in the generating transgenic animals or chromosome and gene mapping, in the generating transgenic animals or chromosome and gene mapping, in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals with genetic disorders as well as for recombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identification. The proteins are useful contracted and proteins are useful in colypeptide (agonists). The polymucleotides and proteins are useful in colypeptide (angonists). The polymucleotides and proteins are useful in colypeptide (angonists). The polymucleotides and proteins are useful in calagnostic assays for PRO eg. detecting its expression in specific colls, itssues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be sequence presented its a PCR primer which was used to amplify a PRO containing and con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; PCR; primer; ss; PRO; secreted; transmembrane; gastrointestinal mucosa; mucosal lesion; skin disease; kertinocyte differentiation; psoriasis; Parkinson's disease; Alstanation; psoriasis; Parkinson's disease; amyotrophic lateral sclerosis; ALS; neuropathy; cell growth; cancer; tumour; viral infection; neurodegenerative disease; antithrombotic agent; haemorrhage; endometrial bleeding angiogenesis; kidney tissue; apoptosis; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nootropic; neuroprotective; cytostatic; virucide; anticoagulant.
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                                                                                                                                                                                                                                                                                                                                                                             polynucleotide of the invention.
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98US-0104080P.

99WO-US021090. 99WO-US021547. 99WO-US023089.

99WO-US020594 99WO-US020944 99WO-US028313. 99WO-US028301. 99WO-US028564.

2000WO-US000219.

99WO-US028214

970S-0059266P 970S-006228F 970S-006348F 970S-006348F 970S-0063144P 970S-0063127P 970S-0063127P 970S-0063127P 970S-0063128P 970S-0064103P 970S-0064168P 970S-006646EP 970S-006646EP 970S-006646EP 970S-006646EP ö

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Score 14.4; DB 1; Length 18; Pred. No. 1.7e+03;

Seguence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

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New PRO polypeptides useful for treating Parkinson's disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal ulceration, Alzheimer's disease, amyotrophic lateral sclerosis and Usher syndrome.
                                                         WPI; 2003-755103/71.
                                                  Williams PM,
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Query Match
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Matches 15; Conservative
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Kljavin IJ;
Tumas D;
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Fong S, Gao W, Gerber H, Gerritsen I
Grimaldi JC, Gurney AL, Hillan KJ, I
an J, Paoni NF, Roy MA, Stewart TA,
22-FEB-2000; 2000WO-US004414.
24-FEB-2000; 2000WO-US005004.
02-MAR-2000; 2000WO-US005841.
20-MAR-2000; 2000WO-US00843.
30-MAR-2000; 2000WO-US00843.
22-MAY-2000; 2000WO-US01644.
02-JUN-2000; 2000WO-US015264.
28-JUN-2000; 2000WO-US021710.
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18-SEP-2000; 2000US-US023328.
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Mather JP, Pan J,
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Filvaroff E,
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Example 36; SEQ ID NO 229; 468pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides have be used to raise and the nucleic acid encoding them. The polypeptides are useful for modulating at least one biological activity of a cell. PRO polypeptide. The PRO plotypeptide are useful for modulating the biological activity of a cell. PRO polypeptide. The PRO polypeptide antibodies ace useful for modulating the biological activity of a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptide. The PRO polypeptide antibodies associated with the preservation and maintenance of gastrointesinal muces and the repair of actual machinal muces and the repair of actual machinal mac

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disease;
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                                                                                                                                  Human secreted/transmembrane protein, #44, PCR primer #2
                                                                                                                                                       Human; PCR; primer; 88; PRO; secreted; transmembrane;
1101 GCTGTCCTCAGGGGAG 1116
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9705-0059117P
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9705-0059266P
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9705-0062285P
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9705-0063286P
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(GETH) GENENTECH INC.

, Goddard A; javin IJ; Ferrara N; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, F Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen M Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, K Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Pan J, Pac. M, Wood WI; Mather JP, F Williams PM,

WPI; 2003-531434/50.

New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO245 or PRO1868, useful in molecular biology, chromosome and gene mapping, in generating antisense RNA and DNA, and in gene therapy.

Example 36; SEQ ID NO 229; 475pp; English.

The Invention discloses isolated known the polypoptides can be used to raise and the nucleic acid encoding them. The polypoptides can be used to raise and the nucleic acid encoding them. The polypoptides for linking a bioactive molecule to a cell expressing a PRO protein and for modulating to bioactive molecule to a cell expressing a PRO protein and for modulating to polypoptides. The PRO polypoptides are useful for modulating the biological activity of a cell expressing a PRO polypoptide. The PRO Dispeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypoptides. PRO polypoptide and an antiterance of acel expressing PRO polypoptides. PRO polypoptide and an antiterance of gastrointestinal muccea and the repair of acute and chronic mucceal differentiation (e.g. poorties) barkinson's disease, Albeimer's differentiation (e.g. poorties), parkinson's disease, Albeimer's also determine and difficularly disease related to uncontrolled call growth, e.g. cancer. PRO polypoptides also serves as tumour specific antigens which may be additionally, disease related to uncontrolled anti-tumour drugs, and are also employed therapeutically in vivo for lessening the effects of viral infection. The PRO polypoptides can be also used in sease; and are also employed therapeutically in vivo for lessening the effects of viral cheparin, in treating other PRO-associated disorders, in modulating cheparin, in treating other PRO-associated disorders, in modulating other properides and their portions effect the expression of genes which have a role in apoptosis. The polymolectides are useful in melecular mapping, in the generating transgenic animals of the forth and protein and for chemographic and protein and pr ; 0 invention discloses isolated PRO secreted/transmembrane polypeptides gastrointestinal mucosa; mucosal lesion; skin disease; keratinocyte differentiation; psoriasis; Parkinson's disease; Alzheimer's diseases; amyotrophic lateral sclerosis; ALS; neuropathy; cell growth; cancer; tumour; viral infection; neurodegenerative disease; Gaps ; 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.76+03; ative 0; Mismatches 1; Indels Human secreted/transmembrane protein, #44, PCR primer #2. Human; PCR; primer; ss; PRO; secreted; transmembrane; Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; 1101 GCTGTCCTCAGGGGAG 1116 ADA42836 standard; DNA; 18 BP. 20-NOV-2003 (first entry) 15; Conservative Best_Local Similarity Matches 15; Conserv ADA42836; Query Match RESULT 1911 ADA42836 à 셤

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antithrombotic agent; haemorrhage; endometrial bleeding angiogenesis; kidney tissue; apoptosis; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nootropic; neuroprotective; cytostatic; virucide; anticoagulant.
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99US-0145698P.
99US-0146222P.
99WO-US020594.
                                 99WO-US020944.
99WO-US021090.
99WO-US021547.
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              99US-0143048P
                                                99WO-US023089
99WO-US028214
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2000WO-US015264.
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24-FEB-2000;
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22-MAY-2000;
                  26-JUL-1999;
28-JUL-1999;
08-SEP-1999;
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20-DEC-1999;
20-DEC-1999;
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                                 13-SEP-1999
                                                     29-NOV-13
                                            15-SEP-
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(GETH) GENENTECH INC.

Botstein D, Desnoyers L, Eaton DL, Ferrara N; Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Grimaldi JC, Gurnes AL, Hillan KJ, Kljavin IJ; an J, Paoni NP, Roy MA, Stewart TA, Tumas D; Wood WI, Ashkenazi A, Botste: Filvaroff E, Fong S Godowski PJ, Grimal Mather JP, Pan J, Williams PM,

WPI; 2003-755052/71.

Novel isolated secreted and transmembrane PRO polypeptide, useful for tissue typing, treating Parkinson's disease, Alzheimer's disease, birth defects, cancer.

Example 36; SEQ ID NO 229; 464pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise at least one biological activity of a cell. PRO polypeptide are useful for detecting other PRO polypeptides in a sample and for linking a for a tleast one biological activity of a cell. PRO polypeptides are useful bloactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. PRO polypeptides are also useful.

CC of acell expressing PRO polypeptides. PRO polypeptides are also useful cor treating disordars associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte and confidence and chronic lateral sclerosis (ALS), neuropathies and diseases, amyotrophic lateral sclerosis (ALS), neuropathies and additionally disease related to uncontrolled cell growth, e.g. cancer. PRO polypeptides also serves as tumour specific antigens which may be exploited as therapeutic targets for anti-tumour drugs, and are also expect as tumour specific antigens which may colypeptides can be also used in assays to determine if it has a role in neurodegenesative diseases or their reversal, as an endometrial bleeding angiogenesis, and may also have an effect on kidney theorem which have a role in apoptosis. The polymucleotides are useful in

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9705-0063128P

9705-0063327P

9705-0063341P

9705-0063542P

9705-0063544P

9705-0063564P

9705-0063564P

9705-0063704P

9705-0066120P

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9705-00666770P

9705-00666772P
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98US-0100262P.
98WO-US019177.
98WO-US019330.
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99WO-US028313
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2000US-00665350
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                                                                        27-0CT-1997;
28-0CT-1997;
28-0CT-1997;
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                  molecular biology including uses as hybridisation probes for CDNA library to isolate the full-length PRO CDNA or to isolate other CDNAs, in chancosome and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides, for generating transgenic animals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals which are useful as probes and for the genetic analysis of individuals with genetic disorders as well as for recombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as the PRO polypeptide (agonists). The polymucleotides and proteins are useful chanced (attaining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific clasmostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from seed in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO entimed the angelian and a second to amplify a PRO entimed the angelian and a series of the angelian and a second to amplify a PRO entimed the angelian and a series and angelian and and and angelian and and and an angelian and and an and and and any and an angelian and and any and an and any a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

0.4%; Score 14.4; DB 1; Length 18;
Best Local Similarity 93.8%; Pred. No. 1.7e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                           polynucleotide of the invention.
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9705-0059117P
9705-0059121P
9705-0059122P
9705-0059263P
9705-0059268P
9705-0062285P
9705-0062285P
9705-0062285P
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17-OCT-1997
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10-JUL-2001; 2001US-00902759.
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The invention relates to human PRO polypeptides (secreted and transmembrane polypeptides) and the polymuclectides encoding them. The polypeptides are useful for detecting PRO polypeptides and for linking a bioactive molecule to a cell expressing the polypeptides, where the bioactive molecule is a toxin, radiolabel or an antibody. The bioactive material causes the death of the cell. The polypeptides or antibodies pecific to the polypeptides are useful for modulating at least one biological activity of a cell expressing the polypeptides. The biological activity of a cell expressing the polypeptides. The biological activity of a cell expressing the polypeptides. The polypeptides are useful for treating disorders associated with leukocyte homing such as asthma, rheumatoid arthritis, psoriasis and multiple clerosis, repair of acute and chronic mucosal lesions such as clerosis, repair of acute and chronic mucosal lesions such as clerosist of the polypeptides. The polymuclectides are useful as hybridization probes, in chromosome and gene mapping, in generation of antisense RNA and DNA, in the preparation of RPCR), Northern analysis, southern analysis and Western analysis. This sequence represents a PCR polymer used in isolation of a human PRO polymuclectide of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gastrointestinal mucosa; mucosal lesion; skin disease; keratinocyte differentiation; psoriasis; Parkinson's disease; Alzheimer's diseases; amyotrophic lateral sclerosis; ALS; neuropathy; cell growth; cancer; tumour; viral infection; neurodegenerative disease; antithrombotic agent; haemorrhage; endometrial bleeding angiogenesis; kidney tissue; apptosis; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nootropic; neuroprotective; cytostatic; virucide; anticoagulant.
                                                                                                                                                                        Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers.
                                            Ferrara N;
n ME, Goddard A;
Kljavin IJ;
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                                            Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Elvaroff E, Pong S, Gao W, Gebber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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ID ADB77755
ID ADB77755
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AC ADB77755;
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DT 04-DEC-2003 (first entry)
XX
XX
DE Human secreted/transmembrane pro
XX
XX
W Human; PCR; primer; 88; PRO; 8ec
XW gastrointestinal mucosa; mucosal
XW Alzheimer's diseases; amyotrophi
XW Alzheimer's diseases; amyotrophi
XW Alzheimer's diseases; amyotrophi
XW call growth, cancer; tumour; vir
XM xidney tissue; apopocosis; therap
XM immunohistochemical staining; ge
XM ycostatic; virucide; anticoagul
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Williams PM, Wood WI;
              (GETH ) GENENTECH INC.
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05-JAN-2000;
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02-JUN-2000; 2000WO-US015264.
28-JUL-2000; 2000WO-US020710.
24-AUG-2000; 2000WO-US023328.
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22-MAY-2000; 2000WO-US014042
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(GETH) GENENTECH INC.

Ferrara N; ME, Goddard A; Kljavin IJ; Tumas D; Eaton DL, F I, Gerritsen N Hillan KJ, F Stewart TA, Ashkenazi A, Botstein D, Desnoyers L, Elivaroff E, Fong S, Gao W, Gerber H, Godowski PJ, Grimaldi JC, Gurney AL, H Mather JP, Pan J, Paoni NP, Roy MA, S Wood WI:

New isolated secreted and transmembrane polypeptide, useful for treating diseases, e.g. Parkinson's disease, Alzheimer's disease, amyotrophic lateral sclerosis, cancer, neuropathies, diabetes and psoriasis.

Example 36; Page 98; 467pp; English

The invention distoless isolated from the polypeptides can be used to raise and the nuclear acid encoding them. The polypeptides can be used to raise antibodies that acid encoding them. The polypeptides can be used to raise at least one biological activity of a cell. PRO polypeptides are useful to deactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide attloades are useful for modulating the bioactive molecule to a cell expressing a PRO polypeptide are also useful configuration and maintenance of a cell expressing PRO polypeptides are proposed are useful for modulating the biological activity of a cell expressing PRO polypeptides are also useful corf gastrointestinal mucosa and the repair of acute and chronic mucosal for treating disorders associated with abnormal keratinocyte molecule (e.g. pooriasis). Parkinson's disease, Alzheimer's diseases, amyotrophic lateral sclerosis (ALS), neuropathies and chronic mucosal clifferentiation (e.g. pooriasis). Parkinson's disease, and the respect of actific antisgens which may be cappetide as therapeutic targets for anti-tumour drugs, and are also additionally, disease related to uncontrolled cell growth, e.g. cancer. CC PRO polypeptides also serves as tumour specific antisgens which may be expression of additionally, disease related to uncontrolled cell growth, e.g. cancer. CC moleculas also serves as tumour specific antisgens which may be expression of antithrombotic agent with reduced risk for haemorrhage as compared with the properties can be also used in assays to determine if it has a role in neurodegenerative diseases or their reversal, as an entit portions affect the expression of concerning the parting other PRO-sasociated disorders, in modulating other PRO-sasociated disorders, in modulating other PRO-sasociated disorders, or modulating of corposeme and gene mapping, in the generation of antisense RNA and DNA, corposed and gene mapping, in the generation relations and the force of sensation and served of the respection and gene mapping, in t The invention discloses isolated PRO secreted/transmembrane polypeptides

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polypeptide (antagonists). The polymucleotides and proteins are useful for tissue typing. PRO antibodies are useful for immunohistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful diagnostic assays for PRO eg. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polymucleotide of the invention.
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gastrointestinal mucosa; mucosal lesion; skin disease;
keratinocyte differentiation; psoriasis; Parkinson's disease;
                                                                                                                                                                              Ouery Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted/transmembrane protein, #44, PCR primer #2
                                                                                                                                                    Seguence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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970S-0059117P.
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970S-0059184P.
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97US-0063120P.
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97US-0063327P.
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97US-0063542P.
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28-OCT-1997;
28-OCT-1997;
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17-OCT-1
21-OCT-1
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05-JAN-2000;
11-FEB-2000;
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28-0CT-1997;
28-0CT-1997;
28-0CT-1997;
29-0CT-1997;
29-0CT-1997;
29-0CT-1997;
29-0CT-1997;
29-0CT-1997;
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16-DEC-1999;
20-DEC-1999;
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31-OCT-1997;
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29-NOV-1999;
30-NOV-1999;
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Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, (GETH) GENENTECH INC. Ashkenazi A, Filvaroff E, Godowski PJ,

Ferrara N; ME, Goddard A; Kljavin IJ;

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodise that specifically bind to the PRO polypeptides for linking a caric dencoding them. The polypeptides can be used to raise antibodise that specifically bind to the PRO polypeptide. For iluking a taleast one biological activity of a cell. PRO polypeptide. The PRO polypeptide are useful for modulating the biological activity of a cell. PRO polypeptide are useful for modulating the biological activity of collaborative molecule to a cell expressing PRO polypeptides. PRO polypeptide are also useful for treating disorders associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, skin diseases associated with abnormal keratinocyte differentiation (e.g. psoriasis), Parkinson's disease, anyotrophic largets for articumour drugs, and are also explosited as therapeutically in vivo for lessening the effects of viral infection. The PRO polypeptides can be also used in assays to determine explosited as therapeutically in vivo for lessening the effects of viral infection. The PRO polypeptides can be also used in assays to determine explosited as role in neurodegenesis, and may also have an effect on kidney tissue. PRO polypeptides can be also used in assays to determine the partin in treating other PRO-associated disorders, in modulating endometrial bleeding and opensors and also have an effect on kidney tissue. PRO polypeptides and their portions affect the expression of successing the protection and their portions and so the construction of antisense RNA and chromosome and gene mapping, in the development and sortening of the processing the protein and for chromosome and protein and for chromosome identification of antisense purpose the represent and for chromosome identification. The PRO polypeptides are useful in the development capents, for screening compounds to identify the protein and for chromos used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynuclectide of the invention. Novel isolated native PRO polypeptide useful for tissue typing, modulating biological activity of cell, as molecular weight markers in protein electrophoresis, for treating enterocolitis, Zollinger-Ellison Paoni NF, Roy MA, Stewart TA, Tumas D; Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Example 36; Page 104; 475pp; English. Mather JP, Pan J, Pac Williams PM, Wood WI; WPI; 2003-765412/72. syndrome.

Gaps ö Query Match

0.4%; Score 14.4; DB 1; Length 18;
Best Local Similarity 93.8%; Pred. No. 1.7e+03;
Matches 15; Conservative 0; Mismatches 1; Indels

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1101 GCTGTCCTCAGGGGAG 1116 3 GCTGTCCACAGGGAG 18 ð g.

ADC28537 standard; DNA; 18 BP RESULT 1915 ADC28537 ID ADC28537 XX

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Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                                                                                           Human secreted/transmembrane protein, #44, PCR primer #2.
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970S-0059113P

970S-0059112P

970S-0059122P

970S-0059122P

970S-0059128P

970S-0062128P

970S-006218P

970S-006281P

970S-006281P

970S-006281P

970S-006381P

970S-006373P

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970S-006378P

970S-006480P

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                                             (first entry)
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29-0CT-1997;
31-0CT-1997;
31-0CT-1997;
03-NOV-1997;
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24-NOV-1997;
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Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating skin, neurodegenerative diseases, as an antithrombotic agent and for inducing endothelial cell apoptosis.
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24-FEB-2000; 2000WO-US005004.

20-MAR-2000; 2000WO-US005841.

20-MAR-2000; 2000WO-US007377.

30-MAR-2000; 2000WO-US008439.

22-MAY-2000; 2000WO-US014042.

02-JUN-2000; 2000WO-US014042.

28-JUL-2000; 2000WO-US015264.
97US-0066772P.
97US-0066840P.
97US-0069425P.
98US-0089803P.
98US-01001824.
98WO-US019177.
98WO-US019310.
98WO-US019477.
98US-0104080P.
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990S-014569BP
990C-014622P-
990C-018020594
990C-018021547
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990C-01802189
990C-018028115
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2000WO-US003565.
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10-SEP-1998;
10-SEP-1998;
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20-DEC-1999;
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05-OCT-1999
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30-NOV-19
01-DEC-19
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08-SEP-1
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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO polypeptide, for linking a taleast one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide are useful as pharmaceuticals, disgnostics, biosensors or polymuclectides are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothelial growth factor (VEGE)-etimulated proliferation of endothelial cells, modulating the proliferation of

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9705-0059184P
9705-005926B
9705-005228B
9705-006228B
9705-00628B
9705-00628B
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9705-0063B
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9705-0066B
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28-0CT-19
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13-OCT-19
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31-0CT-1
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 Stimulated 7-1 ympnocytes, enhanding fine survival or proliferation of cells, modulating glucose or FFA uptake, inducing proliferation and/or rectinal neurons or rod photoreceptor cells, inducing proliferation and/or rectinal membran of chondrocytes. In particular, these are useful for detecting or treating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arbitis) in mammals. PRO polypetides and their portions affect the expression of genes which have a role in cell death. The polymucleotides or arbitis pigmentosum), obligate the full-length PRO cDNA or to isolate other contains and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypetides, for generation of antisense RNA and DNA, for preparing PRO polypetides, for generation of antisense RNA cond DNA, for preparing PRO polypetides, for generation of antisense RNA cond DNA, for preparing PRO polypetides, for generation of antisense RNA cond DNA, for preparing PRO polypetides, for generation of antisense RNA cond DNA, for preparing PRO pulpeptide (alsorders as well as for recombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis or those that mimic the PRO polypeptide (antagonists). The polymucleotides and proteins are useful in diagnostic assays for RNO e.g. detecting its expression in specific cells, tissues or serum and for affinity or expression in specific cells, tissues or serum and for affinity and contibodies are useful in diagnostic assays for RNO e.g. detecting securing a generation gene. The sequence presented is a PCR primer which was used to maplify a PRO polymucleotide of the invention.
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stimulated T-lymphocytes, enhancing the survival or proliferation of
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein, #44, PCR primer #2.
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17-SEP-1997;
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WPI; 2003-540675/51.
        24-AUG-2000;
18-SEP-2000;
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28-OCT-1997
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88888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating skin, neurodegenerative diseases, as an antithrombotic agent and for inducing endothelial cell apoptosis.
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1 ME, Goddard A;
Kljavin IJ;
Tumas D;
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           16-DEC-1999; 99WO-US032031.

20-DEC-1999; 99WO-US03091.

20-DEC-1999; 99WO-US03091.

50-DEC-1999; 99WO-US03091.

11-FEB-2000; 2000WO-US003565.

22-FEB-2000; 2000WO-US00414.

22-FEB-2000; 2000WO-US005841.

20-MAR-2000; 2000WO-US005841.

20-MAR-2000; 2000WO-US008439.

22-MAR-2000; 2000WO-US014042.

22-MAR-2000; 2000WO-US014042.

22-MAR-2000; 2000WO-US014042.

22-MAR-2000; 2000WO-US0120564.
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Filvaroff E, Fong S, Cac
Godowski PJ, Grimaldi JC
"...her JP, Pan J, Paoni
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PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
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                                                                                                                                  Score 14.4; DB 1; Length 18;
Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein, #44, PCR primer #2.
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                                                                                         Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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97US-0059117P.
97US-0059119P.
97US-0059121P.
97US-0059122P.
97US-0059184P.
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9705-0063127P-
9705-0063128P-
9705-0063329P-
9705-0063329P-
9705-0063541P-
9705-0063542P-
9705-0063542P-
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97US-0062125P.
97US-0062285P.
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97US-0062814P.
97US-0062816P.
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97US-0063550P.
97US-0063564P.
97US-0063435P.
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Best Local Similarity 93.8
Matches 15, Conservative
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Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating skin, neurodegenerative diseases, as an antithrombotic agent and for inducing endothelial cell apoptosis.

Example 36; SEQ ID NO 229; 473pp; English

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97US-0063704P.
97US-0063732P.
97US-0063734P.
97US-0063738P.
97US-0063738P.
97US-0063718P.
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97US-0066364P.
97US-0066453P.
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97US-0069425P.
98US-0088026P.
98US-0099803P.
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2000WO-US015264.
2000WO-US020710.
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97US-0065186P.
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97US-0065693P.
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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide for linking a bioactive molecule to a cell expressing a PRO polypeptide for inking a bioactive molecule to a cell expressing a PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide and for linking a coll expressing a PRO polypeptide and activity of a cell expressing a PRO polypeptide or linking a bioactive molecule to a cell expressing a PRO polypeptide or linking a cell expressing PRO polypeptides. The PRO polypeptide or linking a cell expressing PRO polypeptides. The PRO polypeptide or linking a cell expression of a cell expression of proceeders. These are useful as pharmaceuticals, diagnostics, blossnors or collocations. These are useful for stimulating hypertrophy of meantal cells, modulating the proliferation of condorhelial growth factor (VEGF) stimulated proliferation of endochelial cells, inducing cropiferation of condorpoytes, enhancing the proliferation of condorpoytes, inducing proliferation of condorpoytes, inducing proliferation of condorpoytes, inducing cells, modulating cells, modulating cells, modulating proliferation of condorpoytes, inducing proliferation of condorpoytes, inducing proliferation of condorpoytes, inducing the proliferation of condorpoytes, inducing the proliferation of condorpoytes, inducing the proliferation of condorpoytes, or articladed disorders (e.g. sports inducing cells, modulating induced cartical cartical colls, modered, hypoinsulinaemia, or bone or carticaled disorders (e.g. sports inducing cells, hypoinsulinaemia, or bone or carticaled as proteins are useful for conformation of antisens RNA corposation of genes which have a role death. The polymorpoyers are useful in molecular biology including uses as hybridisation of conforted and blay, for preparing RRO polypeptides,

Ferrara N; ME, Goddard A; Kljavin IJ; Tumas D; Desnoyers L, Eaton DL, W, Gerber H, Gerritsen Gurney AL, Hillan KJ, NF, Roy MA, Stewart TA, Filvarur. ... Grimaldı uc, ... Godowski PJ, Grimaldı uc, Mather JP, Pan J, Paoni NF, ... Wood WI; Botstein D, Der Fong S, Gao W, Grimaldi JC, G (GETH) GENENTECH INC. Ashkenazi A, Filvaroff E,

WPI; 2003-540676/51

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antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
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neonatal heart, vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hone disorder; carliage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; annorectic.
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97US-0063542P
97US-0063549P
97US-0063560P
97US-0063564P
97US-0063435P
97US-0063704P
97US-0063732P
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9705-006424BP
9705-0065186P
9705-006564BP
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9705-0066120P
9705-0066120P
9705-006634P
9705-0066453P
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97US-0059117P.
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97US-0059184P.
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97US - 00621259.
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98US-0100262P.
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98US-01008S8P.
98WG-US019437
98US-0109408P.
98WG-US025108.
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99US-0143048P.
99US-014622P.
99WG-US0209944.
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(GETH) GENENTECH INC.

D, Desnoyers L, Eaton DL, Ferrara N; Gao W, Gerber H, Garritsen ME, Goddard A; Goddard A; Hillan KJ, Kljavin IJ; oni NF, Roy MA, Stewart TA, Tumas D; Botstein D, Desno Fong S, Gao W, G Grimaldi JC, Gurn Pan J, Paoni NF, R Mather JP, Pan J, P. William PM, Wood WI; Ashkenazi A, Filvaroff E, Godowski PJ,

WPI; 2003-615762/58.

Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers.

Example 36; SEQ ID NO 229; 476pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for deterting other PRO polypeptides in a sample and for linking a collypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptides or cell expressing a PRO polypeptides or cell expressing a planaceuticals, diagnostics, biosensors or polyncleotides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating hypertrophy of neonatal collecation of endothelial cells, modulating hypertrophy of menatal proliferation of endothelial cells, modulating the proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of cells, modulating proliferation of cells, modulating proliferation and/or receils, modulating or treating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. sports injuries or retinitis pigmentosum), obesity, diabetes, hyperineulianiamia, or bone or cartilage disorders (e.g. sports injuries or

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22-FEB-2000; 2000WO-US004414
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      cc arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polynucleotides are useful in molecular biology including uses as hybridisation probes for conversion probes as the probable of the relation probes conversions. The proparation of antisense RNA can DNA, for preparing PRO polypeptides, for generation of antisense RNA can DNA, for preparing PRO polypeptides, for generation of antisense RNA can DNA, for preparing PRO polypeptides, for generation of antisense RNA can DNA, for preparing the protein are useful in the development and screening of therapeutically useful reagents, as probes and for the combinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polymucleotides and proteins are useful in diagnostic assays for RO e.g. detecting its expression in specific cells, tissues or serum and for affinity can independent of PRO from recombinant cell culture or natural sources. The PRO polypeptide (from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a cepture of amplify a PRO polypurce presented is a PCR primer which was used to amplify a PRO polypurce of the invention.
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99WO-US028313

2000WO-US003565

98WO-US019177. 98WO-US019330. 98US-0100858P. 98WO-US019437.

98US-0104080P. 98US-0109304P. 98WO-US025108.

970S-0062287P 970S-0063486P 970S-0063414P 970S-00631212P 970S-00631212P 970S-0063122P 970S-0063122P 970S-0063128P 970S-0063542P 970S-0063542P 970S-0063542P 970S-0063542P 970S-0063542P 970S-0063544P 970S-0063544P 970S-0063544P 970S-006364P 970S-0063732P 970S-0063732P 970S-0063732P 970S-0063732P 970S-0063734P 970S-0063734P 970S-0063734P 970S-0063734P 970S-0064103P 970S-0064103P 970S-0064103P 970S-006646P 970S-006646P 970S-006646P 970S-006646P 970S-006646P 970S-006646P 970S-006646P 970S-006646P 970S-0066473P 970S-0066473P 970S-0066473P 970S-0066473P 970S-0066473P 970S-0066473P 970S-0066473P

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Score 14.4; DB 1; Length 18; Pred. No. 1.7e+03; Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; 0.4%; Query Match Best Local Similarity

97US-0063870P 97US-0064103P

31-OCT-1997;

Matches 15; Conservative ઠે 셤 The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide for linking a bioactive molecule to a cell expressing a PRO polypeptide for linking a bioactive molecule to a cell expressing a RRO polypeptides. The PRO to detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a RRO polypeptides. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymethodical activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymethodical activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymethodical activity of proliferation of endothelial cells, inducing proliferation of formation of a cartilad activity of proliferation of endothelial cells, inducing proliferation of collar activity of activity of a cells, modulating glucose or FRA uptake, inducing proliferation and/or cells, modulating glucose or FRA uptake, inducing proliferation of chondrocytes. In particular, these are useful for cells, modulating or boseity, diabetes, hyperinalinamia, cancerous cretinitis pigmentosum), obesity, diabetes, hyperinalinamia, cancerous cretinitis pigmentosum), obesity, diabetes, hyperinalinamia, cancerous cretinitis pigmentosum), obesity, diabetes, hyperinalinamia, can be complement or bologopetides and their portion and or sartilage disorders as hybridisation probes or cartilage disorders (e.g. sports injuries or sufficient manners or polymethodical despression of genes which have a role in cell death. The polymethodies are useful in moderally which are useful in the development and screening of therapeutic antendamiantly expressing t purification of PRO from recombinant cell culture or natural sources. The RRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention. Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating cancers, asthma, rheumatoid arthritis, neurological diseases, and skin diseases. ME, Goddard A; Kljavin IJ; Ferrara N; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Isl'Owroff E, Fong S, Gao W, Gerber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Example 36; SEQ ID NO 229; 478pp; English. ; 2000WO-US014042. ; 2000WO-US015264. ; 2000WO-US020710. ; 2000WO-US023328. 2000WO-US007377 2000WO-US008439 18-SEP-2000; 2000US-00665350 Pan J, Paor M, Wood WI;

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                                                                                                                                                                             Human; PCR; primer; 85; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neconatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoroeceptor cell; c-fos; glucose; FRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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                   1101 GCTGTCCTCAGGGGAG 1116
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97US - 0059121P
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Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff B, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Ann Roy M, Stewart TA, Tumas D; Wood WI; Williams PM,

WPI; 2003-585107/55.

Novel isolated PRO polypeptides e.g. PRO234 (useful for treating rheumatoid arthritis, psoriasis and multiple sclerosis) and PRO187 (useful for treating Alzheimer's disease, cancer).

Example 36; SEQ ID NO 229; 451pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise

contibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protesh and for modulating cat it least one biological activity of a cell. PRO polypeptides are useful cat a cell expressing a PRO polypeptides are useful controlled and polypeptides or ports or in polypeptides or ports or including uses as hypridisation probes or polypeptides or polypeptides or ports or polypeptides or polypeptides or ports or polypeptides or ports or polypeptides or pol ö Gaps ö Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; 1101 GCTGTCCTCAGGGGAG 1116

3 GCTGTCCACAGGGAG 18 g

ADC28961 standard; DNA; 18 ADC28961; RESULT 1921

BP

18-DEC-2003 (first entry)

Human secreted/transmembrane protein, #44, PCR primer #2.

tissue typing; immunolistochemical staining; gene therapy;
neonatal heart; vascular endothelial growth factor; VEGF; proliferation;
andothelial cell; stimulated T-lymphocyte; retinal neuron;
rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte;
cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder;
retinitis pigmentosum; obesity; diabete; hyperinaulinaemia;
hypoinsulinaemia; bone disorder; cartilage disorder; sport injury;
arthritis; cardiant; vulnerary; cytostatic; ophthalmological;
osteopathic; antiarthritic; anorectic. Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;

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970S-0059121P.
970S-0059128P.
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970S-0059184P.
970S-0059268P.
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07-JUL-1999;
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05-JAN-2000; 2000WO-US000219.
11-FEB-2000; 2000WO-US00365.
22-FEB-2000; 2000WO-US003661.
24-FEB-2000; 2000WO-US005004.
02-MAR-2000; 2000WO-US00561.
30-MAR-2000; 2000WO-US005841.
31-MAR-2000; 2000WO-US006419. 99WO-US028313. 99WO-US028301. 99WO-US028564. 99WO-US028565. 99WO-US030095. 2000WO-US005841. 2000WO-US007377. 2000WO-US008439. 99WO-US023089 99WO-US028214 99WO-US030911. 02-JUN-2000; 2000WO-US015264. 28-JUL-2000; 2000WO-US020710. 99WO-US020944 99WO-US021090 99WO-US021547 2000WO-US023328 L8-SEP-2000; 2000US-00665350 30-NOV-1999; 01-DEC-1999; 02-DEC-1999; 16-DEC-1999; 16-DEC-1999; 20-DEC-1999; 28-JUL-1999; 08-SEP-1999; 13-SEP-1999; 15-SEP-1999; 15-SEP-1999; 05-OCT-1999;

(GETH) GENENTECH INC.

Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; Williams PM, Wood WI;

WPI; 2003-615797/58.

Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating skin, neurodegenerative diseases, as an antithrombotic agent and for inducing endothelial cell apoptosis.

Example 36; SEQ ID NO 229; 470pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the mucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO polypeptide are useful for detecting other PRO polypeptides in a sample and for modulating a bloactive molecule to a cell expressing a PRO polypeptide are useful for bloactive molecule to a cell expressing a PRO polypeptide are useful for polypeptide are useful for scimulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide are useful as pharmaceuticals, diagnostics, blosensors or comply and are useful as pharmaceuticals, diagnostics, blosensors or bioreactors. These are useful for stimulating hypertrophy of neonatal complying vascular endothelial growth factor (VEGF)-stimulated complying vascular endothelial cells, modulating the proliferation of complying vascular endothelial cells, modulating co-fos in endothelial cells, modulating proliferation of chondrocytes, endoting proliferation and chondrocytes, inducing co-fos in endothelial cells, modulating prosecore properties of sight due to cells, modulating glucose or FPA uptake, inducing proliferation and chondrocytes. In particular, these are useful for reatinal disorders or injuries (e.g. ports injuries or retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymolymolecular polymolecular molecular molecular molecular molecular molecular molecular molecular and gene mapping, in the generation of and second molecular properting second molecular properting properting properting second molecu

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genetic analysis of individuals with genetic disorders as well as for the combinantly expressing the protein and for chromsome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polymucleotides and proteins are useful for tissue typing. PRO antibodies are useful for immunohistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, itssues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
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ADC40846
ID ADC40846
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AC ADC40846
XX
DT 18-DC-
XX
DE Human;
XX
Human;
XW
Human;
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ADC19503 standard; DNA; 18 BP.

RESULT 1923 ADC19503 (first entry)

18-DEC-2003

ADC19503;

(GETH) GENENTECH INC.

Ferrara N; ME, Goddard A; Kljavin IJ; Tumas D; Desnoyers L, Eaton DL, W, Gerber H, Gerritsen Gurney AL, Hillan KJ, NF, Roy MA, Stewart TA, Botstein D, Desnoyers L, Fong S, Gao W, Gerber H Grimaldi JC, Gurney AL, an J, Paoni NF, Roy MA, Wood WJ; Filvaror-Godowski PJ, Grima. TP. Pan J, Mather JP, Pa Williams PM, Ashkenazi A, Filvaroff E,

WPI; 2003-708343/67.

Novel PRO polypeptides useful for treating Parkinson's disease, Alzheimer's disease, enterocolitis, Zollinger-Ellison syndrome, psoriasis, epidermoid carcinoma of the vulva and gliomas, gynecological diseases.

Example 36; SEQ ID NO 229; 473pp; English.

The invention discloses isolated who secreted/transmembrane polypeptides on the mucleic acid encoding them. The polypeptides can be used to raise and the mucleic acid encoding them. The polypeptides can be used to raise bloactive molecule ace acell expressing a PRO protein and for inking a call expressing a PRO polypeptide. The PRO polypeptides are useful acivity of a cell. PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptide. The PRO polypeptide antibodies are useful for stimulating hypertrophy of neonatal beart, inhibiting vascular endothelial growth factor (VEGP) stimulated controllers are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothelial growth factor (VEGP) stimulated controllers of endothelial cells, modulating proliferation of endothelial cells, modulating proliferation of endotrocytes. In particular, these are useful for etimal neurons or rod photoreceptor cells, inducing proliferation and/or receival modulating glucose or FRA uptake, inducing proliferation and/or receival modulating glucose or FRA uptake, inducing proliferation and/or receival modulating or treating candar inagification of formatics (e.g. loss of sight due to retinities priments, retinial disorders or injuries (e.g. loss of sight due to retinities) in momentals, properties and their portions affect the expression of genes which have a role in cell death. The polymolecotides for cumular or knockout animals with genetic and pub. for preparing PRO polypeptides and their portions are useful in molecular marker for protein and for the genetic analysis of individuals with genetic disorders as well as therapeutic agents, for percenting compounds to identify the genetic analysis of individuals with genetic process, as therapeutic agents, in the generation of entering or therapeutic analysis of individuals wit antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention. The invention discloses isolated PRO secreted/transmembrane polypeptides

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels 93.8%; Best Local Similarity 93.8 Matches 15, Conservative Query Match

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1101 GCTGTCCTCAGGGGAG 1116 GCTGTCCACAGGGGAG 18

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tissue typing; immunolistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                                                                transmembrane; therapeutic;
                                              Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                 PCR; primer; ss; PRO; secreted;
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9705 - 0063045P
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97US-0062814P.
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22-MAY-2000; 2000WO-US014042
02-JUN-2000; 2000WO-US015264
28-JUL-2000; 2000WO-US015264
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18-SEP-2000; 2000US-00665350
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22-FEB-2000; 2000WO-US004414
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02-MAR-2000; 2000WO-US005841
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William PM, Wood WI;
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Ferrara N; 1 ME, Goddard A; Kljavin IJ; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Filvaroff E, Fong S, Gao W, Gebber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Novel isolated PRO polypeptide useful for treating Parkinson's disease, enterocolitis, Zollinger-Ellison syndrome, gastrointestinal ulceration, Alzheimer's disease, amyotrophic lateral sclerosis.

Example 36; SEQ ID NO 229; 478pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or

registration of endothelial cells, modulating the proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of endothelial cells, modulating proliferation of cells, modulating proliferation of cells, modulating proliferation and/or receiptor cells, inducing c-fos in endothelial cells, modulating glucose or FFA uptake, inducing c-fos in endothelial cells, modulating glucose or FFA uptake, inducing c-fos in endothelial cells, modulating glucose or FFA uptake, inducing c-fos in endothelial detecting or treating cardiac insufficiency disorders, wounds, cancerous cumours, retinal disorders or injuries (e.g. loss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides are useful in molecular biology including uses as hybridisation probes or companies, in chromosome and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides, for generating transgenic cond by, for preparing PRO polypeptides, for generating transgenic and DNA, for preparing PRO polypeptides, as probes and for the generating of therapeutically useful reagents, as probes and for the generating are useful sampled in the development and generation are useful as molecular marker for protein electrophoresis proteins are useful for tissue typing; PRO antibodies are useful for the PRO polypeptide (antagonists). The polymucleotides and proteins are useful for tissue typing. PRO antibodies are useful for the pRO polypeptide (antagonists) or prevent the effect of immunohistochemical staining and/or assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity and produce presented is a PRR primer which was used to amplify a PRO polymerice. polynucleotides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothelial growth factor (VEGF)-stimulated

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ; 0 Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels

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RESULT 1924

ADC33951 standard; DNA; 18 BP ADC33951;

18-DEC-2003 (first entry)

Human secreted/transmembrane protein, #44, PCR primer #2.

tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF, proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; hypoinsulinaemia; bone disorder; cartilage disorder; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritis; anorectic. Human, PCR, primer, ss, PRO, secreted, transmembrane, therapeutic;

Homo sapiens.

US2003073077-A1

17-APR-2003

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97US-0059113P.
97US-0059115P.
97US-0059117P.
97US-0059119P.
97US-0059121P.
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24-FEB-2000; 2000WO-US005004. 02-MAR-2000; 2000WO-US005841. 20-MAR-2000; 2000WO-US005437. 30-MAR-2000; 2000WO-US004377. 22-MAY-2000; 2000WO-US014042. 02-UJN-2000; 2000WO-US01564. 28-UJL-2000; 2000WO-US02328. 99WO-US028313. 99WO-US028301. 99WO-US028564. 99WO-US028565. 99WO-US03095. 99WO-US030999 2000WO-US000219 2000WO-US003565. 2000WO-US004414. 18-SEP-2000; 2000US-00665350 05-JAN-2000; 2 11-FEB-2000; 2 24-FEB-2000; 2 02-MAR-2000; 2 00-MAR-2000; 2 30-MAR-2000; 2 22-MAY-2000; 2 02-JUN-2000; 2 20-DEC-1999;

(GETH) GENENTECH INC.

Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NP, Roy MA, Stewart TA, Tumas D; Godowski PJ, Grimaldi Mather JP, Pan J, Pac Williams PM, Wood WI;

WPI; 2003-695953/66.

Novel isolated PRO polypeptides e.g. PRO245 and PRO1868, useful for treating e.g. Parkinson's disease, Alzheimer's disease, amyotrophic lateral sclerosis, cancer, neuropathies, diabetes and psoriasis.

Example 36; SEQ ID NO 229; 476pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides can the mucleic acid encoding them. The polypeptides can be used to raise cantibodies that specifically bind to the PRO polypeptides can be useful continuing a bioactive molecule to a call expressing a PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a call expressing a PRO polypeptides are useful for detecting other PRO polypeptides. The PRO polypeptides or polypeptide are useful for modulating the biological activity of a call expressing as PRO polypeptides or polypeptides are useful for modulating the biological activity of a call expressing PRO polypeptides. The PRO polypeptides or polymentication for a call expressing the properties or polymucleotides are useful for modulating the proliferation of candential calls, modulating the proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of candential calls, inducing proliferation of stimulated T-lymphocytes, inputake, inducing c-fost in endothelial calls, modulating glucose or FFA uptake, inducing proliferation of calls, modulating glucose or FFA uptake, inducing proliferation of calls, modulating glucose or FFA uptake, inducing proliferation of cardiac insufficiency disorders, womans, or bone or cartilage disorders, womans, or bone or cartilage disorders, possibly dispersed by modulating properties in molecular properties and thair portions affect the arthritis pigmentosum), obesity diabetes, hyperinsulinamia, or bone or cartilage disorders or carthritis or consone and genes which have a role in call dearth. The polymocal and planty to isolate the full-length PRO cDNA or to isolate other constants in molecular biology including uses as hybridisation care useful is molecular marker for protein electronaly of therapeutically useful reagents, as probes and for the generic analysis of individuals with generic disorders of artisp

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970S-0063732P
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99WO-US028301
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Mather JP, P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-FEB-2000;
24-FEB-2000;
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Filvaroff E,
28-0CT-1997,
28-0CT-1997,
29-0CT-1997,
29-0CT-1997,
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29-0CT-1997,
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31-OCT-1997;
03-NOV-1997;
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25-NOV-19
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18-NOV-19
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21-NOV-19
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               antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity urification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; oradiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                                                                                                                                                                                                    Gaps
   immunohistochemical staining and/or assay of sample fluids. Anti-PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic;
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                                                                                                                                                                0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                   Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                   1101 GCTGTCCTCAGGGGAG 1116
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97US-0059117P.
97US-0059117P.
97US-0059121P.
97US-0059122P.
97US-0059263P.
97US-0059268P.
97US-0062125P.
97US-0062125P.
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97US-0063045P.
97US-0063120P.
97US-0063121P.
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                                                                                                                                                                                                                                                                GCTGTCCACAGGGGAG 18
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                                                                                                                                                                                                  15; Conservative
                                                                                                                                                                                   Local Similarity
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17-SEP-1997;
17-SEP-1997;
17-SEP-1997;
18-SEP-1997;
15-OCT-1997;
17-OCT-1997;
24-OCT-1997;
24-OCT-1997;
24-OCT-1997;
24-OCT-1997;
24-OCT-1997;
24-OCT-1997;
24-OCT-1997;
24-OCT-1997;
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27-0CT-1997;
28-0CT-1997;
28-0CT-1997;
28-0CT-1997;
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                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                 RESULT 1925
ADC13021
XX
AC ADC13022
XX
XX
DT 18-DEC-
XX
XX
DE Human 8
XX
Cardiac
XW endothe
XW endothe
XW cardiac
XW cardiac
XW ACONO
XX
CARDIA
XX
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18-SEP-2000; Z000US-00655350.
(GETH) GENENTECH INC.
Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;

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contribudies that specifically bind to the PRO puppetides can be used to raise antibodies that specifically bind to the PRO polypeptides, for linking a can biological activity of a cell. PRO polypeptides for linking a can biological activity of a cell. PRO polypeptides are useful bioactive molecule to a cell expressing a PRO polypeptide. The RO polypeptides are useful for modulating the biological activity of bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptides are useful for modulating the biological activity of placeture molecule are useful for modulating the biological activity of placeture. These are useful for simulating hypertrophy of meomatal bioacative molecules are useful for simulating hypertrophy of meomatal conformation of endothelial cells, modulating the proliferation of placeture are useful for simulating propertrophy of meomatal neart, inhibiting vascular endothelial cells, inducing c-fos in endothelial cells, modulating inducing c-fos in endothelial cells, modulating induced or FRA uptake, inducing c-fos in endothelial cells, modulating proliferation of the central nearnon of chondrocytes. In particular, these are useful for cetinal nearnon of chondrocytes, in particular, these are useful for cetinal nearnon of chondrocytes. In particular, these are useful for cetinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, chors or reatinged cardiac insufficienty disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymocell conductions are useful in molecular biology including uses as hybridisation probes for cDNA ibrary to isolate the full-length pro conducts and the development and conductions and gene mapping, in the generation of antisons conducting of therapeutically useful reagents, or special sarbing of therapeutically useful reagents, or processing the protein and entities are useful and propertied charagonists for genetic analysis of individuals with g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                            Novel isolated secreted and transmembrane PRO polypeptides e.g. PRO245 and PRO1868, useful for treating e.g. Parkinson's disease, Alzheimer's disease, amyotropic lateral sclerosis, cancer, neuropathies, diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
                                                                                                                                                                                                                                                                invention discloses isolated PRO secreted/transmembrane polypeptides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                              Example 36; SEQ ID NO 229; 473pp; English.
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Wood WI
                                               WPI; 2003-743809/70.
Williams PM,
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ADC12473 standard; DNA; 18 BP.
3 gérérécacacacada 18
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                                                                                      ADC12473;
                                    RESULT 1926
                                                ADC12473
ID ADC1
XX ADC1
AC ADC1
XX I8-D
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97US-0066840P.

25-NOV-1997; 12-DEC-1997;

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tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fcs; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinaulinaemia; hypoinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; one disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                           transmembrane; therapeutic;
         Human secreted/transmembrane protein, #44, PCR primer #2.
                            primer; ss; PRO; secreted;
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9705-005912P-
9705-0059184P-
9705-0059263P-
9705-0059266P-
9705-0062285P-
9705-0062285P-
9705-0062287P-
9705-0062287P-
9705-0062887P-
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970S-0063045P-
970S-0063121P-
970S-0063128P-
970S-0063128P-
970S-0063329P-
970S-0063542P-
970S-0063542P-
970S-0063542P-
970S-0063542P-
970S-0063543P-
970S-0063734P-
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970S-00637338P-
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970S-0064218P-
970S-0064218P-
970S-0064248P-
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97US-0066120P.
97US-0066364P.
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                                                                                                                                                          US2003082541-A1.
                                                                                                                                       Homo sapiens.
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15-OCT-1997;
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03-NOV-1
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28-OCT-
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24-0CT-
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98US-0088026P.
98US-0099803P.
98WO-US018824.
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                              98WO-US019330
                                    98US-0100858P
                                           98WO-US019437
                                                98US-0104080P
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30-MAR-2000;
22-MAY-2000;
                 14-SEP-1998;
14-SEP-1998;
16-SEP-1998;
17-SEP-1998;
17-SEP-1998;
13-OCT-1998;
20-NOV-1998;
                                                            01-DEC-1998;
22-DEC-1998;
07-JUL-1999;
26-JUL-1999;
28-JUL-1999;
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08-SEP-1999;
13-SEP-1999;
15-SEP-1999;
15-SEP-1999;
05-OCT-1999;
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01-DEC-1999;
02-DEC-1999;
02-DEC-1999;
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(GETH) GENENTECH INC.

Ferrara N; 1 ME, Goddard A; Kljavin IJ; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NP, Roy MA, Stewart TA, Wood WI: Williams PM,

WPI; 2003-743881/70.

New secreted transmembrane PRO polypeptides and nucleic acids encoding the polypeptides, useful in gene therapy, in identifying chromosomes, as chromosome markers, in generating probes and in tissue typing.

Example 36; SEQ ID NO 229; 487pp; English.

and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide for linking a biactorie molecule to a cell expressing a PRO polypeptide. for linking at least one biological activity of a cell. PRO polypeptides are useful to detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptide or polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptide or polymetrophy of neonatal chart, inhibiting vascular endothelial growth factor (VEGP)-stimulated proliferation of endothelial cells, modulating the proliferation of crimulated T-lymphocytes, enhancing the survival or proliferation of retinal neurons or rod photoreceptor cells, inducing c-fos in endothelial cells, modulating proliferation and/or re The invention discloses isolated PRO secreted/transmembrane polypeptides

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cumoniary are treating cardiac insufficiency disorders, wounds, cancerous detecting or treating cardiac insufficiency disorders, wounds, cancerous cumours, retinal disorders or injuries (e.g. 10ss of sight due to retinitis pigmentosum), obesity, diabetee, hyperingulinaemia, hyperingulinaemia, or bone or cartilage disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a note in cell death. The polymuclectides are useful in molecular biology including uses as hybridisation probes of or cDNA library to isolate the full-length RNO cDNA or to isolate other CDNAs, in chromosome and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides, for generating transgenic animals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the corrections are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify those there is the minimal which protein and for offromosome identification. The proteins are useful as molecular marker for protein electrophoresis or purposes, as therapeutic agents, for screening compounds to identify to those that minim the RNO polypeptide (ancagonists). The polymucleotides and proteins are useful for tissue typing. PRO antibodies are useful for antibodies are useful for managonists). The polymucleotides and proteins are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity continued or matural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polypurcleotide of the invention.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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1101 GCTGTCCTCAGGGGAG 1116 Query Match
Best Local Similarity 93.8%; 3 ocrerccacacacas 18 셤

RESULT 1927 ADD05028

ADD05028 standard; DNA; 18 01-JAN-2004 (first entry) ADD05028;

Human secreted/transmembrane protein, #44, PCR primer #2.

tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsullinaemia; hypoinsullinaemia; hone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic. Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;

Homo sapiens.

US2003104469-A1.

17-JUL-2001; 2001US-00907652. 17-SEP-1997; 05-JUN-2003.

97US-0059113P. 97US-0059115P. 97US-0059117P. 97US-0059119P. 97US-0059121P. 17-SEP-1997; 17-SEP-1997; 17-SEP-1997; 17-SEP-1997;

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97US-0059184P.
97US-0059263P.
97US-0059266P.
                            97US-0062125P.
97US-0062285P.
97US-0062287P.
97US-0063486P.
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98US-0099803P.
98WO-US018824.
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97US-0064809P.
97US-0065186P.
                                                          97US-0062814P.
97US-0062816P.
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12-NOV-1997;
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2000MO-US005641. 2000MO-US007377. 2000MO-US00439. 2000MO-US016264. 2000MO-US015564. 2000MO-US023328. 99WO-US030999. 2000WO-US00319. 2000WO-US003565. 2000WO-US004414. 2000WO-US005004. 2000US-00665350 11-FEB-2000; 22-FEB-2000; 24-FEB-2000; 02-MAR-2000; 20-MAR-2000; 30-MAR-2000; 22-MAY-2000; 24-AUG-2000; 18-SEP-2000;

(GETH) GENENTECH INC.

Filvaroff B. Forstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff B, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D: Williams PM, Wood WI;

WPI; 2003-801231/75.

Novel isolated native PRO polypeptide useful for tissue typing, modulating biological activity of cell, as molecular weight markers in protein electrophoresis, for treating enterocolitis, Zollinger-Ellison syndrome.

Example 36; SEQ ID NO 229; 474pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nurolic acid encoding them. The polypeptides can be used to raise and the nurolic acid encoding them. The polypeptide to raise antibodies that specifically bind to the PRO polypeptide. For linking a control system wolecule to a cell systessing a PRO polypeptide. The PRO polypeptide are useful for a cell systessing a PRO polypeptide on the properties of the prop

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Novel isolated native PRO polypeptide useful for treating Parkinson's disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal
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1 ME, Goddard A;
Kljavin IJ;
Tumas D;
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97US-0063734P
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97US-0065846P
97US-0065634P
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98WO-US025108.
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21-NOV-1
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                                                                                                                                                                                                                                                                                                    tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; oardiant; vulnezary; cytostatic; ophthalmological; osteopathic; antiarthritis; anorectic.
                                                                            Gaps
                                                                                                                                                                                                                                                                                           Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;
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                                                   0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03;
                                                                            1; Indels
                                                                                                                                                                                                                                                                   Human secreted/transmembrane protein, #44, PCR primer #2.
                          Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
  amplify a PRO polynucleotide of the invention.
                                                                          0; Mismatches
                                                                                                    1101 GCTGTCCTCAGGGGAG 1116
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97US - 0059117P
97US - 0059121P
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9705-0063128-
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                                            Ouery Match
Best Local Similarity 93.8
Matches 15; Conservative
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28-0CT-1997;
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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides con the used to raise and the nucleic acid encoding them. The polypeptide for linking a carticular polypeptide of the used to raise the least one biological activity of a cell. PRO polypeptide. The PRO polypeptide activity of a cell expressing a PRO polypeptide. The PRO polypeptide activity of a cell perpension and for linking a to load tive molecule to a cell expressing a PRO polypeptides. The PRO polypeptide activity of polypeptide antibodies are useful for modulating the biological activity of polypeptide antibodies are useful for modulating the biological activity of polypeptide activity of a cell expressing PRO polypeptides. The PRO polypeptides of collpany of collpa
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tissue typing; immunohistochemical staining; gene therapy;
neonatal heart; vascular endothelial growth factor; VEGF; proliferation;
ulceration, Alzheimer's disease, amyotrophic lateral sclerosis, Usher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted/transmembrane protein, #44, PCR primer #2.
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                                                               Example 36; SEQ ID NO 229; 487pp; English
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endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diserte: hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological;
                                                  osteopathic; antiarthritic; anorectic
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                                                                    Homo sapiens.
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                      98WO-US025108
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20-DEC-1999;
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29-NOV-1999
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Ferrara N; n ME, Goddard A; Kljavin IJ; Desnoyers L, Eaton DL, F W, Gerber H, Gerritsen N Gurney AL, Hillan KJ, F NF, Roy MA, Stewart TA, Paoni NF, Botstein D, Des Fong S, Gao W, Grimaldi JC, Gu Filvaror. Godowski PJ, Gr., Ther JP, Pan J, Pau.. Ashkenazi A, Filvaroff E,

WPI; 2003-801268/75.

Novel isolated native PRO polypeptide useful for tissue typing, modulating biological activity of cell, as molecular weight markers in protein electrophoresis, for treating enterocolitis, Zollinger-Ellison

Example 36; SEQ ID NO 229; 472pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating a to reast one biological activity of a cell. PRO polypeptides are useful compactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide are useful compactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptides or polypeptide antibodies are useful for modulating the biological activity of a cell expressing a pRO polypeptides or coll expressing PRO polypeptides or polymucleotides are useful for stimulating hypertrophy of neonatal colliferation of the factor (VEGP) stimulated colliferation of endothelial growth factor (VEGP) stimulated colliferation of conditions or red photoreceptor cells, inducing colliferation of chondrocytes, enhancing the survival or proliferation of conditions or cells, inducing colliferation of chondrocytes, inducing proliferation and/or reciplementation of chondrocytes, inducing proliferation and conditions cells, modulating proliferation of chondrocytes, inducing proliferation of chondrocytes, inducing colliferation of chondrocytes, inducing colliferation of chondrocytes, inducing colliferation of chondrocytes, inducing colliferation of chondrocytes, inducing proliferation of chondrocytes, inducing colliferation colliferation of chondrocytes, chondrocytes, chondrocytes, chondr

cc arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polynucleotides care useful in molecular biology including uses as hybridisation probes for cDNA library to isolate the full-length PRO cDNA or to isolate other contained to preparing PRO polypeptides, for generation of antisense RNA and DNA, for preparing PRO polypeptides, for generation of antisense RNA and DNA, for preparing languagemic canimals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the crecombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoreas purposes, as therapeutic agents, for screening compounds to identify the proteins are useful as molecular marker for protein electrophoreas purposes, as therapeutic agents, for screening compounds to identify the PRO polypeptide (antagonists). The polynuciectides and proteins are useful for tissue typing PRO antibodies are useful for commission-marker and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity corresponded to the sequence presented is a PCR primer which was used to explicate amplify a PRO polymucleotide of the invention. Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö Score 14.4; DB 1; Length 18; Pred. No. 1.7e+03; 0; Mismatches 1; Indels 1101 GCTGTCCTCAGGGGAG 1116 0.4%; Query Match
Best Local Similarity 93.8
Matches 15; Conservative ઠે

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RESULT 1930

3 GCTGTCCACAGGGGAG 18

ADD44211 standard; DNA; 18 ADD44211;

ВЪ.

15-JAN-2004 (first entry)

Carboxypeptidase G2 (CPG2) enzyme mutagenic oligonucleotide OL576.

bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; BC 3.4.17.11; mutagenic; ss

Pseudomonas sp. RS-16. Synthetic

WO2003045426-A1.

05-JUN-2003

27-NOV-2002; 2002WO-EP013351

29-NOV-2001; 2001EP-00128519. 25-JAN-2002; 2002EP-00001778. 13-SEP-2002; 2002EP-00020634.

(MERE) MERCK PATENT GMBH.

Carr FJ; ŝ Hellendoorn K, Baker M, Williams

WPI; 2003-513617/48.

New modified bacterial enzyme carboxypeptidase G2 (CPG2) having substantially non-immunogenic or less immunogenic than any non-modified CPG2, useful for inducing an immune response in a human host.

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ADE34862;
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                                                            carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any on-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations compared with the non-modified parochial enzyme. The alterations cause a reduction or an elimination of one or more of T-cell captops sequences, which act in the parental enzyme as MHC class II binding ligands and stimulate T-cells. The modified CPG2 enzyme and the CPG2 proteins have immunostimulant activity and may be used in enzyme therapy. The modified CPG2 enzyme may be used to induce an immune response in a human host, or as a therapeutic entity such as the gene directed enzyme prodrug strategy. The peptide is useful for the manufacture of a modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo.

This polymodelectide sequence represents a mutagenic oligonuclectide used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            directed enzyme prodrug strategy; vaccine; enzyme; EC 3.4.17.11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels
                                               invention relates to a novel modified bacterial enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in the production of a modified CPG2 gene of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 9 A; 2 C; 4 G; 3 T; 0 U; 0 Other;
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4; Page 37; 52pp; English
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25-JAN-2002; 2002EP-00001778.
13-SEP-2002; 2002EP-00020634.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15; Conservative
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ADD44227/c
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The invention relates to a novel modified bacterial enzyme car result in CPG2 carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations cause a reduction or an elimination of one or more of T-cell epitope sequences, which act in the parental enzyme as MHC class II epitope sequences, which act in the parental enzyme as MHC class II epitope sequences, which act in the parental enzyme as MHC class II epitope sequences, which act in the parental enzyme as II epitope sequences, which act in the parental enzyme as MHC class II epitope sequences, which act in the parental enzyme as MHC class II epitope sequences, which act in the parental enzyme as II enzyme therapy. The modified CPG2 enzyme may be used in enzyme therapy. The modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo.

This polymucleotide sequence represents a mutagenic objectide used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in the production of a modified CPG2 gene of the invention.
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97US-0059117P.
97US-0059117P.
97US-0059121P.
97US-0059124P.
97US-0059263P.
97US-0059266P.
97US-0062285P.
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97US-0062814P.
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Best Local Similarity 93.8°
Matches 15, Conservative
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17-OCT-1997;
21-OCT-1997;
24-OCT-1997;
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97US-0063121P.
97US-0063127P.
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24-FEB-2000; 2000WO-US005004
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30-MAR-2000;
22-MAY-2000;
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13 - SEP - 1999;
15 - SEP - 1999;
05 - OCT - 1999;
30 - NOV - 1999;
01 - DEC - 1999;
02 - DEC - 1999;
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24-0CT-1997;
24-0CT-1997;
27-0CT-1997;
28-0CT-1997;
28-0CT-1997;
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28-OCT-1997;
28-OCT-1997;
28-OCT-1997;
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29-OCT-1997;
29-OCT-1997;
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24-NOV-1997;
24-NOV-1997;
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16-SEP-1998;
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17-NOV-1997
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26-NOV-1997;
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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide. for linking a caribodies that specifically bind to the PRO polypeptide. for linking a transmission molecule to a cell expressing a PRO polypeptide. The PRO polypeptide is to reach a cell expressing a PRO polypeptide in the PRO polypeptide antibodies are useful for modulating the biological activity of a cell. PRO polypeptide on the properties of a cell expressing PRO polypeptides. The PRO polypeptides or polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymolectides are useful for modulating operation of properties of a cell expressing properties of the properties of polymolectides are useful for setuminating propertopy of meanaral chart. inhibiting varcular endothelial growth factor (VEGP) estimulated proliferation of condorcoves. In particular, inducing creating cardiacling glucose or FRA uptack, inducing proliferation of chondrocyes. Cells, modulating proliferation of chondrocyes. In particular, these are useful for cells, modulating proliferation of chondrocyes. In particular, these are useful for detecting cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to continuing or treating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the arthritis in mammals, proposed or cartilage disorders and phylication of sense which have a rote in cell for coNA library to isolate the full-inger Research of antisense RNA continuing or knockout animals which are useful in the development and continuing of therapeutically useful responders as probes and for the properties of individuals with generic disorders as well as molecular manary crecentin New isolated PRO polypeptides e.g. PRO245 and PRO1868, useful for treating e.g. Parkinson's disease, Alzheimer's disease, amyotrophic lateral sclerosis, cancer, neuropathies, diabetes and psoriasis. Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; Williams PM, Wood WI; Example 36; SEQ ID NO 229; 474pp; English. 02-JUN-2000; 2000WO-US015264. 28-JUL-2000; 2000WO-US02710. 24-AUG-2000; 2000WO-US023328. 18-SEP-2000; 2000US-00665350. (GETH) GENENTECH INC. WPI; 2003-777194/73.

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match

0.4%; Score 14.4; DB 1; Length 18;
Best Local Similarity 93.8%; Pred. No. 1.7e+03;
Matches 15; Conservative 0; Mismatches 1; Indels

ö

1101 GCTGTCCTCAGGGGAG 1116

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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated PRO polypeptides e.g. PRO211, PRO217 and PRO230, useful for treating Parkinson's disease, Alzheimer's disease, amyotropic lateral sclerosis, cancer, neuropathies and psoriasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Botstein D, Desnoyers L, Eaton DL, Ferrara N; , Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; wood WI;
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         97US-0066120P.
97US-0066451P.
97US-0066451P.
97US-0066410P.
97US-0066770P.
97US-006640P.
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99WO-US028313.
99WO-US028301.
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2000WO-US005004.
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2000WO-US023328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GETH ) GENENTECH INC.
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02-DEC-1999;
20-DEC-1999;
20-DEC-1999;
20-DEC-1999;
05-JAN-2009;
05-JAN-2000;
11-FEB-2000;
24-FEB-2000;
24-FEB-2000;
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20-MAR-2000;
30-MAR-2000;
22-MAY-2000;
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Filvaroff E,
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Williams PM,
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                  21-NOV-1997;
24-NOV-1997;
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24-NOV-1997;
24-NOV-1997;
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13-OCT-19
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01-DEC-19
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Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; tissue typing; immunohistochemical staining; gene therapy; encontal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; daabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                                                                                                                    Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                 9705-0059113P
9705-0059113P
9705-0059111P
9705-0059121P
9705-0059124P
9705-0059128P
9705-005926F
9705-006228F
9705-006228F
9705-0062814P
9705-0062814P
9705-0063121P
9705-006354P
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9705-006373B.
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9705-006424BP.
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9705-006424BP.
                                                     BP,
GCTGTCCACAGGGGAG 18
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                                                     ADH59345 standard; DNA; 18
                                                                                               (first entry)
                                                                                                                                                                                                                                                                                   US2003039972-A1
                                                                                                                                                                                                                                                                Homo sapiens.
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21-0CT-1997;
24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
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27-0CT-1997;
28-0CT-1997;
28-0CT-1997;
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17-SEP-1997;
17-SEP-1997;
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17-SEP-1997
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29-OCT-1997
                                                                          ADH59345
                               RESULT 1933
                                           ADH59345
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20-MAR-2003

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bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide are useful as remained for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or cell expressing PRO polypeptides. The PRO polypeptides or polymucleotides are useful for ethmilating hypertrophy of meomatal heart, inhibiting vascular endethelial growth factor (VGGP)-etimulated colls. These are useful for ethmilating propertrophy of neomatal heart, inhibiting vascular endethelial cells, modulating the proliferation of crimulated cells, modulating the proliferation of crimulated cells, modulating the proliferation of crimulated reliance. The second of the condrocytes. In particular, these are useful for cells, modulating quicose or PRA witake, inducing proliferation of chondrocytes. In particular, these are useful for detecting or treating cardiac innelficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to rathitials) in mammals. PRO polypeptides and thair portions affect the arthitial in molecular biology including uses as hybridisation probes or carthitials in molecular biology including uses as hybridisation probes or constant of an endomental process and gene wasping, in the generation of antisense RNA and DNA, for preparing RNO polypeptides, for generating transgenic and surface or commands or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals with genetic disorders as well as for creening of therapeutically useful reagents, as probes and for the proteins are useful in diagnostic assay of sample fluids. Anti-PRO these that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (agonists) or prevent the PRO polypeptide call in a sequence presented in specific cells, tissues popularic assay of sample f
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Gaps ö 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.78+03; tive 0; Mismatches 1; Indels Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; 1101 GCTGTCCTCAGGGGAG 1116 Query Match
Best Local Similarity 93.8
Matches 15; Conservative

GCTGTCCACAGGGGAG 18 ઠ g

ADI38124 standard; DNA; 18 BP ADI38124;

22-APR-2004 (first entry)

tissue typing; immunohistochemical staining; gene therapy;
neonatal heart; vascular endothelial growth factor; VEGF; proliferation;
endothelial cell; stimulated T-lymphocyte; retinal neuron;
rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte;
cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder;
retinitis pigmentosum; obesity; diabete; hyperinaulinaemia;
hypoinsulinaemia; bone disorder; cartilage disorder; sport injury;
arthritis; cardiant; vulnerary; cytostatic; ophthalmological;
osteopathic; antiarthritic; anorectic. Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; Human secreted/transmembrane protein, #44, PCR primer #2.

Homo sapiens

US2003054352-A1

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9705-0059115P

9705-0059117P

9705-005912P

9705-005926P

9705-0059268P

9705-0059268P

9705-006285P

9705-0062814P

9705-0062814P

9705-0062814P

9705-0063121P

9705-0063121P

9705-0063121P

9705-0063121P

9705-0063121P

9705-0063124P

9705-0063124P

9705-0063124P

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9705-0063124P

9705-0063124P

9705-0063128P

9705-0066468P
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98US-0088026P.
98US-0099803P.
                   17-JUL-2001; 2001US-00907925
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98WO-US019177
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24-FEB-2000; 2000WO-US005004.

20-MAR-2000; 2000WO-US005841.

20-MAR-2000; 2000WO-US008439.

22-MAY-2000; 2000WO-US014042.

02-JUN-2000; 2000WO-US015264.

28-JUL-2000; 2000WO-US015264.
                                          0095.
                                              99WO-US030911.
                                                                    2000WO-US004414.
                          99WO-US028301
                                                         2000WO-US000219
2000WO-US003565
                                                                                                             24-AUG-2000; 2000WO-US023328.
18-SEP-2000; 2000US-00665350.
          99WO-US023089
                99WO-US028214
                     99WO-US028313
                                99WO-US028564
                                    99WO-US028565
                                          99WO-US03
                                                         05-JAN-2000; 2
11-FEB-2000; 2
22-FEB-2000; 2
24-FEB-2000; 3
                    30-NOV-1999;
01-DEC-1999;
02-DEC-1999;
               29-NOV-1999
                                    02-DEC-1999
16-DEC-1999
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                                                    20-DEC-1999
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Ä, Ferrara N; ME, Goddard A Kljavin IJ; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Williams PM, Wood WI;

WPI; 2003-695899/66.

Novel isolated native PRO polypeptide useful for treating Parkinson's disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal ulceration, Alzheimer's disease, amyotrophic lateral sclerosis, Usher syndrome.

Example 36; SEQ ID NO 229; 471pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise cation the propertied of the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO polypeptides are useful for detecting other PRO polypeptides are useful for detecting other pRO polypeptides are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or coll expressing PRO polypeptides or collection of collection of endothalial cells, modulating the proliferation of collection of endothalial cells, modulating the proliferation of collection of endothalial cells, modulating proliferation of collecting or rod photoreceptor cells, inducing proliferation and/or recollecting or treating cardiac innegration of strainal neurons or rod photoreceptor cells, inducing proliferation of collecting or treating cardiac innegration of collecting or treating cardiac innegration of collecting or treating cardiac innegration of strainal disorders or injuries (e.g. ioso of sight due to collecting or treating cardiac innegration or collecting or collecting or collecting cardiac innegration or collecting cardiac innegration or collecting and their portions affect the collecting or preparing PRO polypeptides and their portions affect the collecting or collecting properties and their portions of antisense RNA and bNA, for preparing PRO polypeptides, for generating transgenic combasion of generic analysis of individuals with generation of combasion. The proteins are useful in the development and gene mapping, in the generation of combas

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purposes, as therapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polynucleotides and proteins are useful for tissue typing. PRO antibodies are useful for immunobistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; PRO; primer; ss; secreted polypeptide; transmembrane polypeptide; pathological disorder; cardiac insufficiency disorder; protein secretion; pancreas; diabetes; gastrointestinal mucosal mucosal lesion; psoriasis; skin disease; keratinocyte differentiation; epithelial cancer; tumour; lung squamous cell carcinoma; epidermoid carcinoma; vulva; glioma; PCR; cytostatic; cardiant; endocrine; antidiabetic; gastrointestinal;
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                                                                                                                                                                                                                 Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                     Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiulcer; dermatological; vulnerary
                                                                                                                                                                                                                                                                               1101 GCTGTCCTCAGGGGAG 1116
                                                                                                                                                                                                                                                                                                                                                                                    ACA59084 standard; DNA; 18 BP.
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9708-0059117P
9708-0059121P
9708-0059124P
9708-0059184P
9708-0059263P
9708-0059268P
9708-0062285P
9708-006228FP
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97US-0063541P.
97US-0063542P.
                                                                                                                                                                                                                                                                                                        GCTGTCCACAGGGGAG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO PCR primer #96.
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Also polypeptides and polynucleotides can be used in treating pathological disorders and tumours, in therapeutic treatment of cardiac insufficiency disorders and in therapeutic treatment of disorders involving protein secretion by the pancreas, including diabetes. They can also be used in treating disorders associated with the preservation and maintenance of gastrointestinal mucosa and the repair of acute and chronic mucosal lesions, and skin diseases associated with abnormal keratinocyte differentiation (e.g., psoriases, epithelial cancers such as lung squamous cell carcinoma, epidermoid carcinoma of the vulva and gliomas). The sequences can be used as molecular markers for protein electrophoresis purposes and can be utilised in protein-protein binding assays, biochemical screening assays, immunoassays and cell-based assays. polynucleotide of the invention
                                                                                                                                                                    Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                      1101 GCTGTCCTCAGGGGAG 1116
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Best Local Similarity 93.8
Matches 15; Conservative
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  97US-0063544P.
97US-0063569P.
97US-0063564P.
97US-0063435P.
97US-0063732P.
97US-0063732P.
97US-0063738P.
97US-0063738P.
97US-0063738P.
97US-0063738P.
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97US-0065186P-
97US-0065186P-
97US-006593P-
97US-0066430P-
97US-0066453P-
97US-0066466P-
97US-0066466P-
97US-0066461P-
97US-0066471P-
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98WO-US018824.
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02-MAR-2000; 2000WO-US005841.
20-MAR-2000; 2000WO-US007377.
30-MAR-2000; 2000WO-US0084397.
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22-FEB-2000; 2000WO-US004414
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  28-OCT-1997;
28-OCT-1997;
28-OCT-1997;
29-OCT-1997;
29-OCT-1997;
29-OCT-1997;
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31-OCT-1997;
31-OCT-1997;
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16-SEP-1998;
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Gaps

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Score 14.4; DB 1; Length 18; Pred. No. 1.7e+03; 0; Mismatches 1; Indels

93.88;

3 gérérécacadadada 18

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PCR primer #106 used to isolate cDNA encoding a human PRO polypeptide.
                                                                                Human; secreted and transmembrane protein; PRO polypeptide; cancer; Alzheimer's disease; ischaemia; cytostatic; nootropic; vasotropic; neuroprotective; PCR; primer; ss.
                                                                                                                                                                                           9705-0059115P
9705-0059117P
9705-0059121P
9705-0059124P
9705-0059184P
9705-0062185P
9705-0062185P
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9705-0063186P
9705-0063187P
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9705-0063121P
9705-0063121P
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9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
                BP
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                ACAS8481 standard; DNA; 18
                                                 10-JUN-2003 (first entry)
                                                                                                                                   US2002192659-A1
                                                                                                                    Homo sapiens.
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17-0CT-1997;
24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
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                                                                                                                                                    19-DEC-2002.
                                                                                                                                                                                               17-SEP-1997
                                 ACA58481;
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RESULT 1936
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Isolated nucleic acid useful for e.g., treating pathological disorders encodes a secreted or transmembrane protein.

Ferrara N; 1 ME, Goddard A; Kljavin IJ; Tumas D;

Desnoyers L, Eaton DL, W, Gerber H, Gerritsen Gurney AL, Hillan KJ, NF, Roy MA, Stewart TA,

Gurney AL, IF, Roy MA,

Paoni NF,

Pan J,

Wood WI;

Williams PM,

Mather JP,

WPI; 2003-328338/31.

Ashkenazi A, Botstein D, Der Filvaroff E, Fong S, Gao W, Godowski PJ, Grimaldi JC, G

The invention relates to human PRO polypeptides (secreted or transmembrane polypeptides) and the polynucleotides encoding them. The

Example 37; Page 102; 473pp; English.

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970S-0065846P-
970S-0065839P-
970S-0066120P-
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Ferrara N;
1 ME, Goddard A;
Kljavin IJ;
Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Elvaroff E, Fong S, Gao W, Gerbber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Wood WI; Mather JP, P

WPI; 2003-361832/34.

New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO245 or PRO1868, useful in molecular biology, chromosome and gene mapping, in generating antisense RNA and DNA, and in gene therapy.

Example 37; Page 102; 474pp; English.

The present invention relates to the isolation of novel human secreted and transmembrane proteins (PRO polypeptides), and the polynucleotide

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sequences encoding them. The polynucleotide sequences are useful in molecular biology, as hybridisation probes, in chromosome and gene mapping, in generating antisense RNA and DNA, and in gene therapy. The polynucleotide sequences may also be used in preparing PRO polypeptides by recombinant techniques, and in generating either transgenic animals or knock-out animals which, in turn, are useful in the development and screening of therapeutically useful reagents. The PRO polypeptides or their antibodies are useful in preparing a medicament for treating condition responsive to the polypeptide or antibody, such as cancer, allahelmer's disease or ischaemia, and in various diagnostic assays. The present sequence represents a PCR primer used in the examples of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinaulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide. For linking a card the modified to a cell expressing a PRO polypeptide. The PRO polypeptide is a sample and for modilating a bloactive molecule to a cell expressing a PRO polypeptides or useful for detecting other PRO polypeptides in a sample and for linking a bloactive molecule to a cell expressing a PRO polypeptides or linking a conscience of a cell expressing PRO polypeptides. The PRO polypeptides or polymolectides are useful for modulating the biological activity of control procedures. These are useful for modulating the proliferation of polymolectides are useful for seimulating phycetrophy of meonatal control procedures. The propose of PRO polypeptides or training a glucose or FRA uptack, inducing proliferation of chondrocytes. Entantacing the proliferation of chondrocytes. In particular, these are useful for ceils, modulating glucose or FRA uptack, inducing proliferation of chondrocytes. In particular, these are useful for ceils, modulating glucose or FRA uptack, inducing proliferation and or the proliferation of chondrocytes. In particular, these are useful for ceils, modulating plucose or FRO uptacks, inducing proliferation of chondrocytes or injuries (e.g. ions of sight due to ceils, modulating plucose or FRO uptacks or injuries (e.g. sport or ceils, modulating plucose or farminase (e.g. sport or ceins of genes which have a role in cell or cord library to isolate the fer ull-length RRO colypeptides and their portions affect the creaming of genes which have a role in cell for cord library to isolate the fer ull-length RRO colypeptides or control or solate or the control or solate the coll or cord library to isolate the fer ull-length RRO polypeptide, or cord library to isolate the fer ull coll or cord library to isolate the fer ull coll or cord library to solate the properties of solate the PRO polypep Novel isolated native PRO polypeptide useful for tissue typing, modulating biological activity of cell, as molecular weight markers in protein electrophoresis, for treating enterocolitis, Zollinger-Ellison Gaps Botstein D, Desnoyers L, Baton DL, Ferrara N; Fong S, Gao W, Gerber H, Gerriteen ME, Goddard A; Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; an J, Paoni NF, Roy MA, Stewart TA, Tumas D; ;; 0 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Example 36; SEQ ID NO 229; 483pp; English. 1101 GCTGTCCTCAGGGGAG 1116 GCTGTCCACAGGGGAG 18 ADE79307 standard; DNA; 18 Ashkenazi A, Bocc.
Filvaroff E, Fong S, Goc Godowski PJ, Grimaldi JC.
Ther JP, Pan J, Paoni Query Match Best Local Similarity 93.81 Matches 15; Conservative WPI; 2003-708341/67. syndrome. RESULT 1938 ADE79307 ID ADE79307 8 56

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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO polypeptide, for linking a test one biological activity of a cell. PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polypeptides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating hypertrophy of meonatal heart, inhibiting vascular endothelial growth factor (VEGF)-stimulated
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tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VBGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; oardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                                                                              Human, PCR, primer, 88, PRO, secreted, transmembrane, therapeutic;
                                                       Human secreted/transmembrane protein, #44, PCR primer #2.
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Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; Williams PM, Wood WI;
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20-MAR-2000; 2000WO-US007377.
30-MAR-2000; 2000WO-US008439.
22-MAY-2000; 2000WO-US014042.
02-UJN-2000; 2000WO-US015284.
28-JUL-2000; 2000WO-US015284.
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22-FEB-2000; 2
24-FEB-2000; 2
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                                                                   10-SEP-1998;
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20-DEC-1999;
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9705-0059117P

9705-0059113P

9705-0059121P

9705-0059264P

9705-0059266P

9705-0062287P

9705-0062184P

9705-0062125P

9705-0062114P

9705-0063144P

9705-00631121P

9705-0063121P

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9705-0063124P

9705-006354P

9705-006354P
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9705-0064103P
9705-006448P9
9705-0065186P
9705-0065618P
9705-006663P
9705-006646P
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98US-0100262P.
98WO-US019177.
98WO-US019330.
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                                                                                                                                                          17-0CT-1997;
21-0CT-1997;
24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
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01-DEC-1999;
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24-0CT-1
24-0CT-1
27-0CT-1
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           proliferation of endothelial cells, modulating the proliferation of etimulated T-lymphocytes, enhancing the survival or proliferation of cetimal neurons or rod photoreceptor cells, inducing c-fos in endothelial cells, modulating glucose or FFA uptake, inducing proliferation and/or redifferentiation of chondrocytes. In particular, these are useful for detecting or treating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides are useful in molecular biology including uses as hybridisation probes for cDNAs, in chromosome and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides, for generating transgenic animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals with genetic disorders as well as for complainantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify the protein are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify the PRO polypeptide (angonists). The polymucleotides and proteins are useful for tissue typing. PRO antibodies are useful for tissue typing. PRO antibodies are useful for useful for tissue typing. PRO antibodies ample fluids. Anti-PRO antibodies and protein are useful for tissue typing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                     antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; one disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1101 GCTGTCCTCAGGGGAG 1116
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tes 15; Conservative
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PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

88988

0.4%;

Query Match 0.4: Best Local Similarity 93.8 Matches 15; Conservative

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Gaps

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Score 14.4; DB 1; Length 18; Pred. No. 1.7e+03; 0; Mismatches 1; Indels

1101 GCTGTCCTCAGGGGAG 1116

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3 gererecacadedas 18

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Ferrara N;
1 ME, Goddard A;
Kljavin IJ;
Tumas D;
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Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen I
Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, I
Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Williams PM, Wood WI;
                                                                                                      2000WO-US000219.
2000WO-US003565.
2000WO-US004414.
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2000WO-US007377.
2000WO-US008439.
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                                                             99WO-US030911.
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                                            99WO-US030095
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                                                                                                                                                                                                                                                                                                                                                                            (GETH ) GENENTECH INC.
02-DEC-1999;
02-DBC-1999;
16-DEC-1999;
20-DBC-1999;
20-DBC-1999;
16-FBB-2000;
22-FBB-2000;
22-FBB-2000;
24-FBB-2000;
20-MAR-2000;
30-MAR-2000;
32-MAR-2000;
22-MAR-2000;
                                                                                                                                                                                                                                                                                           28-JUL-2000;
                                                                                                                                                                                                                                                                                                                                      L8-SEP-2000;
                                                                                                                                                                                                                                                                                                                   24 - AUG - 2000;
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New PRO nucleic acid, useful for manufacturing a medicament for diagnosing or treating tumor or for tissue typing. WPI; 2004-020353/02.

Example 36; SEQ ID NO 229; 480pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the mucleic acid encoding them. The polypeptides contribution and the mucleic acid encoding them. The polypeptides, for linking a bloative molecule to a cell expressing a PRO polypeptide, for linking a bloative molecule to a cell expressing a PRO polypeptides are useful activity of a cell. PRO polypeptides are useful other PRO polypeptides in a sample and for inking a bloactive molecule to a cell expressing a PRO polypeptides are useful other PRO polypeptides. The PRO polypeptide arthodises are useful as pharmaceuticals, diagnostics, biosensors or polymucleotides are useful for stimulating the biological activity of a cell expressing as pharmaceuticals, diagnostics, biosensors or polymucleotides are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothalial cells, modulating the proliferation of proliferation of endothalial cells, modulating the proliferation of stimulated Trymphosytes, enhancing the survival or proliferation of chondrocytes, enhancing the survival or proliferation of chondrocytes, enhancing the survival or proliferation of chondrocytes, enhancing the proliferation of chondrocytes, enhancing proliferation of chondrocytes, enhancing proliferation of stimulated to treating or treating cort reading carder in particular, these are useful for cells, modulating glucose or FFA uprake, inducing proliferation and/or recent cells, modulating of processor or injuries (e.g. loss of sight due to architis pigmentosum), obseity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or retinitis pigmentosum), obseity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or construction in molecular biology including uses as hybridiate one cartilage disorders (e.g. sports injuries or construction in molecular biology including uses as hybridiates on the construction of encoded and public proparating PRO polypeptides, for generating transgenic animals or

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tissue typing; immunohistochemical staining; gene therapy. Lustapeutlic; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological;
                                                                                                     Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic;
                                                                                 Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                                                                                   osteopathic, antiarthritic; anorectic
                                                                                                                                                                                                                                                                                                         970S-0059113P.
970S-0059113P.
970S-0059113P.
970S-0059121P.
970S-0059121P.
970S-0059124P.
970S-0059265P.
970S-006285P.
970S-006285P.
970S-0062814P.
970S-0062816P.
970S-0063120P.
970S-0063121P.
970S-0063121P.
970S-0063121P.
                    ADE73407 standard; DNA; 18 BP
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                                                             29-JAN-2004 (first entry)
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                                                                                                                                                                                                                         Homo sapiens.
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                                         ADE73407;
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27-OCT-1
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RESULT 1940
           ADE73407
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vivlemore401-10.rng

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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic arid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide. For linking a cariobdise that specifically bind to the PRO polypeptide. For linking a to discative molecule to a cell syrpessing a PRO polypeptide. The PRO polypeptides are useful for modulating the biological activity of a cell. PRO polypeptides are useful for modulating the biological activity of a cell. PRO polypeptides on the processing PRO polypeptides. The PRO polypeptides on the processing PRO polypeptides. The PRO polypeptides on the processing PRO polypeptides in a sample and for individual process. These are useful for modulating propertion, of conducting the proliferation of polymeropy of meancal care that individual spluces or FAY uptake, inducing proliferation of conducytes, enhancing the proliferation of conducytes, enhancing the proliferation of conducytes, enhancing the proliferation of conducytes, inducing proliferation of conducytes, enhancing the proliferation of conducytes, inducing proliferation of conducytes, inducing proliferation of conducytes or individual proliferation of denancy or cartiling ediocyters (e.g. sports individual proliferation of solate the proliferation of solate the conduction of antibution of solate the bill-length PRO toby pertices of conduction properties of conduction of solate the full-length PRO toby antibution of therapeutically useful reagents as probes and for the protein and proliferation of the protein and proliferation of the properties of individuals with generation of solate the PRO polypeptide, conducy in solate the PRO poly
New nucleic acids encoding polypeptides designated PRO have sequence identity to various secreted proteins and transmembrane proteins and are useful in molecular techniques and as therapeutic agents.
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0.4%; Score 14.4; DB 1; Length 18;
Best Local Similarity 93.8%; Pred. No. 1.7e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tissue typing; immunohistochemical staining; gene therapy;
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                                                                              Example 36; SEQ ID NO 229; 474pp; English
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n ME, Goddard A;
Kljavin IJ;
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2000WO-US014042.
2000WO-US015264.
2000WO-US020710.
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Mather JP, Pan J, Pac
Williams PM, Wood WI;
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05-JAN-2000;
11-FEB-2000;
22-FEB-2000;
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                                  29-OCT-1997;
29-OCT-1997;
29-OCT-1997;
29-OCT-1997;
31-OCT-1997;
31-OCT-1997;
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20-MAR-2000;
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15-SEP-1999;
15-SEP-1999;
05-OCT-1999;
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18-NOV-1997
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02-DEC-1999
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Gaps

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neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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9705-0059113P

9705-0059113P

9705-0059121P

9705-0059122P

9705-0059128P

9705-0059268P

9705-0062818P

9705-0062818P

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97US-0064215P.
97US-0063870P.
97US-0064103P.
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97US-0064809P.
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97US-0065693P.
97US-0066120P.
97US-0066364P.
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97US-0066466P.
97US-0066511P.
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97US-0066840P.
97US-0069425P.
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98US-0100262P.
98WO-US019177.
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                                                                                                                     US2003148370-A1.
                                                                                                Homo sapiens.
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17-0CT-1997;
17-0CT-1997;
24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
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24-0CT-1997;
27-0CT-1997;
27-0CT-1997;
28-0CT-1997;
28-0CT-1997;
28-0CT-1997;
28-0CT-1997;
28-0CT-1997;
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29-0CT-1997;
29-0CT-1997;
29-0CT-1997;
29-0CT-1997;
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10-SEP-1998;
14-SEP-1998;
14-SEP-1998;
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17-SEP-1997;
17-SEP-1997;
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31-OCT-1997;
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07-NOV-1997;
12-NOV-1997;
17-NOV-1997;
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21-NOV-1997;
21-NOV-1997;
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24-NOV-1997
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98US-0100858P.
98WO-US019437.
98US-0109408P.
98WO-US025108.
99US-0113296P.
99US-0145698P.
99WG-0146222P.
99WG-US020994.
                                                                       99WO-US028313.
99WO-US028301.
99WO-US028564.
99WO-US028565.
99WO-US030095.
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2000WO-US005841.
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2000WO-US015264.
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                                                                                          16-DEC-1999;
20-DEC-1999;
20-DEC-1999;
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24-FEB-2000;
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                 20-NOV-1998;
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Ferrara N; 1 ME, Goddard A; Kljavin IJ; Tumas D; Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NF, Roy MA, Stewart TA, Mather JP, Pan J, Pa Williams PM, Wood WI; Ashkenazi A, Filvaroff E, Godowski PJ,

WPI; 2004-020440/02.

Isolated secreted and transmembrane PRO nucleic acids and the proteins they encode, e.g. PRO245, PRO269 and PRO1868, useful for preventing, diagnosing and treating e.g. disorders relating to blood coagulation.

Example 36; SEQ ID NO 229; 1pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detering other PRO polypeptides in a sample and for linking a collypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides or the PRO polypeptides are useful for modulating the biological activity of a cell expressing PRO polypeptides or the PRO polypeptides are useful for stimulating the biological activity of a cell expressing PRO polypeptides or the PRO polypeptides or proliferation of endothelial cells, modulating the proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of cells, modulating proliferation of cells, inducing collecting acadiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to thypoinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or

WPI; 2004-021580/02.

cc arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polynucleotides care useful in molecular biology including uses as hybridisation probes for cDNA library to isolate the full-length PRO cDNA or to isolate other cDNAs, in chromosome and gene mapping, in the generation of antisense RNA cand DNA, for preparing PRO polypeptides, for generation of antisense RNA cand DNA, for preparing PRO polypeptides, for generation of antisense RNA canimals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the crecombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polynucleotides and proteins are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity cantibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity or profice may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polymucleotide of the invention. Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; tive 0; Mismatches 1; Indels 1101 GCTGTCCTCAGGGAG 1116 Best Local Similarity 93.8 Matches 15; Conservative Query Match 8

Gaps

; 0

3 ecrerceacadeaaa 18

ADE99496 standard; DNA; 18 BP 12-FEB-2004 (first entry) ADE99496;

Human secreted/transmembrane protein, #44, PCR primer #2.

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neconatal heart; vascular endothelial growth factor; VBGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuro; rod photoreceptor cell; c-fos; glucose; FRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartiage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritis; anorectic. RESULT 1942
ADE99496
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AC ADE99496
XX
DT 12-FEB-2
XX
DT 12-FEB-2
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CADE90496
CADE904

Homo sapiens.

JS2003211576-A1.

13-NOV-2003.

18-NOV-2002; 2002US-0029893.

22-FEB-2000; 2000WO-US004414. 18-SEP-2000; 2000US-00665350.

GETH) GENENTECH INC.

, Botstein D, Desnoyers L, Eaton DL, Ferrara N; , Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; , Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; , Wood WI; Ashkenazi A, Filvaroff E, Godowski PJ, Mather JP, E Williams PM,

contribudies that specifically bind to the RPO bypeptide, for linking a chitchodies that specifically bind to the RPO polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise bioactive modecule to a cell expressing a RPO polypeptides. The RPO polypeptides in a sample and for linking a for detecting other RPO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a RPO polypeptide. The PRO polypeptides are useful for modulating the biological activity of a cell expressing a RPO polypeptides. The PRO polypeptides or polymorlectides are useful for simulating hypeptide or control for simulating hypertcophy of mematal bioactors. These are useful for simulating hypertcophy of mematal bioactors. These are useful for simulating propertcophy of mematal control feration of endothelial cells, modulating proliferation of condroverse or FA uptake, inducing or proliferation of condroverse in particular, these are useful for citimal neurons or rod photoreceptor cells, inducing proliferation and/or rectinal neurons or rod photoreceptor cells, inducing proliferation and/or redetering or treating cardiac insufficiently, these are useful for chemory testinal disorders or injurias (e.g. loss of sight due to certinitis pigmentosum), obesity, diabetes, hyperinsulinsemia, or control of genes which have a role in cell death. The polymorlectides are useful in molecular biology including uses as hybridisation probes for cDNA inbrary to isolate the full-length RPO cDNA or to isolate other containing of therapeutically useful respension of antisense RNA care useful in molecular biology including uses as hybridisation of antisense containally expression and gene mapping, in the generation of antisense containing of therapeutically useful respension of genes which have a role in cell death. The polymorphic combinantly expressing the protein and for cure or carting of therapeutically useful respense, as probes and for the growing and properse, as therapeutically u The invention discloses isolated PRO secreted/transmembrane polypeptides New PRO polypeptide for preparing a medicament for treating a condition that is responsive to the PRO polypeptide or anti-PRO antibody, e.g. inflammatory diseases, cancer or acquired immunodeficiency syndrome. Example 36; SEQ ID NO 229; 476pp; English

PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention. Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö Score 14.4; DB 1; Length 18; Pred. No. 1.7e+03; 0; Mismatches 1; Indels 93.8%; Query Match 0.4 Best Local Similarity 93.8 Matches 15, Conservative

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1101 GCTGTCCTCAGGGGAG 1116 3 GCTGTCCACAGGGGAG 18 ò

ADE98615 standard; DNA; 18 ADE98615; ADE98615

RESULT 1943

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Human secreted/transmembrane protein, #44, PCR primer #2. 12-FEB-2004 (first entry)

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tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinaulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnezary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
         Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;
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97US - 0063121P
97US - 0063123P
97US - 0063327P
97US - 0063341P
97US - 0063341P
97US - 0063344P
97US - 0063544P
97US - 0063564P
97US - 0063564P
97US - 0063732P
97US - 0063734P
97US - 0063738P
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97US-0059117P.
97US-0059117P.
97US-0059121P.
97US-0059124P.
97US-0059184P.
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97US-0064809P.
97US-0065186P.
97US-0065846P.
97US-0065633P.
97US-0066120P.
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97US-0062816P.
97US-0063045P.
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97US-006641P.
97US-0066511P.
97US-0066770P.
97US-006840P.
97US-006942F.
98US-0099803P.
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97US-0062285P.
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                                                                                                                      Homo sapiens.
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17-SEP-1997;
17-SEP-1997;
17-SEP-1997;
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24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
24-0CT-1997;
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17-OCT-1997;
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28-OCT-1997;
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10-SEP-1998;
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21-OCT-1997
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29-OCT-1997,
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24-NOV-1997
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21-NOV-1
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98US-0100858P. 98WO-US019437. 98US-0104080P. 98US-0103304P. 98US-0113296P. 99US-0113296P. 99US-0143048P. 98US-0100262P. 98WO-US019177. 98WO-US019330. 99WO-US020594. 99WO-US021090. 99WO-US028301. 99WO-US030095. 2000WO-US004414. 99WO-US021547 99WO-US023089 99WO-US028214 99WO-US028565 99WO-US030999 99WO-US028313 18-SEP-2000; 2000US-00665350 11-FEB-2000; 22-FEB-2000; 24-FEB-2000; 02-MAR-2000; 20-MAR-2000; 20-NOV-1998; 20-DEC-1999; 29-NOV-1 02-DEC-1

(GETH) GENENTECH INC.

Botstein D, Desnoyers L, Eaton DL, Ferrara N; Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; an J, Paoni NF, Roy MA, Stewart TA, Tumas D; Wood WI; Ashkenazi A, Botste Filvaroff E, Fong S Godowski PJ, Grimal Mather JP, Pan J, Williams PM,

WPI; 2004-021576/02.

New isolated native PRO polypeptide useful for treating Parkinson's disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal ulceration, Alzheimer's disease, amyotrophic lateral sclerosis, or Usher

Example 36; SEQ ID NO 229; 469pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for modulating a bloactive molecule to a cell expressing a PRO polypeptide are useful to polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or collocations are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating hypertrophy of neonatal chart, inhibiting vascular endothelial growth factor (VEGF)-stimulated proliferation of endothelial cells, modulating the proliferation of endothelial cells, inducing c-fos in endothelial cells, modulating glucose or FRA uptake, inducing proliferation and/or redifferentiation of chondrocytes. In particular, these are useful for cifferential or cells, modulating proliferation and/or redifferentiation of chondrocytes. In particular, these are useful for

cc detecting or treating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arbititis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides or are useful in molecular biology including uses as hybridisation probes or constant of genes which have a role in cell death. The polymucleotides or are useful in molecular biology including uses as hybridisation probes or constant of preparing PRO polypeptides. For generation of antisense RNA and DNA, for preparing PRO polypeptides, for generation of antisense RNA conformation of therapeutically useful reagents, as probes and for the screening of therapeutically useful reagents, as probes and for the combinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify the proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify the proteins are useful and/or assay of sample fluids. Anti-protein electing its constitution for tissue typing. PRO antibodies are useful for chromosome identification of PRO form recombinant cell culture or natural sources. The profice may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to maplify a PRO polypurcleotide of the invention.

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; iive 0; Mismatches 1; Indels Local Similarity 93.8 hes 15; Conservative Query Match Matches

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ADE99042 standard; DNA; 18 BP ADE99042

ADE99042;

(first entry) 12-FEB-2004 Human secreted/transmembrane protein, #44, PCR primer #2.

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; PRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinaulineamia; hypoinsulinaemia; bone disorder; cartiage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.

Homo sapiens

JS2003211568-A1.

13-NOV-2003.

12-JUL-2001; 2001US-00904805.

16-SEP-1998; 98WO-US019330. 22-FEB-2000; 2000WO-US004414. 97US-0063327P 18-SEP-2000; 2000US-00665350 27-OCT-1997;

(GETH) GENENTECH INC

Ferrara N; ME, Goddard A; Kljavin IJ; Tumas D; Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NP, Roy MA, Stewart TA, Pan J, Pao M. Wood WI; Ashkenazi A, Filvaroff E, Godowski PJ, Williams PM, Mather JP,

WPI; 2004-021575/02.

New secreted and transmembrane nucleic acids and polypeptides, designated as PRO, useful for treating inflammation, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, or

Example 36; SEQ ID NO 229; 473pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the mucleic acid enceding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide for linking a bloactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide in the properties of a cell expressing a PRO polypeptide in the PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptide in the PRO polypeptide antibodies are useful for modulating the biological activity of polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymorlocides are useful for stimulating hypertrophy of neonatal bioreactors. These are useful for stimulating hypertrophy of neonatal proliferation of endothelial cells, modulating the proliferation of crimulate or rod phortoreceptor cells, inducing c-fos in endothelial cells modulating glucose or PRA uptake, inducing proliferation of crimulating glucose or PRA uptake, inducing proliferation and/or recills modulating glucose or PRA uptake, inducing proliferation and/or recills modulating of chondrocytes. In particular these are useful for retinalisemals, or bone or cartilage disorders, wounds, cancerous crimulates properson of genetical and plantary of socretical and plantary and plantary of socretical and plantary and p amplify a PRO polynucleotide of the invention.

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels 93.8%; Query Match 0.4 Best Local Similarity 93.8 Matches 15, Conservative

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Gaps

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RESULT 1945 ADG40512

(WOOD/) WOOD W I. 34-OCT-1997; (GODD/) (MATH/) typing. THE STATE OF THE S

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ADG40512 standard; DNA; 18 BP
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ADG40512;

26-FEB-2004 (first entry)

Human secreted/transmembrane protein, #44, PCR primer #2.

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nemonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsullinaemia; hypoinsullinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.

Homo sapiens.

JS2003225253-A1.

04-DEC-2003.

29-MAY-2003; 2003US-00448923

16-SEP-1998; 98WO-US019330. 30-NOV-1999; 99WO-US028313. 22-PEB-2000; 2000WO-US04414. 18-SEP-2000; 2000US-00665350. 12-JUL-2001; 2001US-00905125.

GODDARD A. GODOWSKI P J. DESNOYERS L. DESN/)

GURNEY A L. MATHER J P. WILLIAMS P M. GURN/)

Goddard A, Godowski PJ, Gurney AL, Mather JP; Wood WI; Desnoyers L, Williams PM,

WPI; 2004-022084/02.

New PRO nucleic acid, useful for manufacturing a medicament for diagnosing or treating tumor, for chromosome mapping or for tissue

Example 36; SEQ ID NO 229; 463pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bloactive molecule to a cell expressing a PRO polypeptide are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide artibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymedicotides are useful for stimulating hypertrophy of neonatal chart, inhibiting vascular endothelial growth factor (VEGF)-stimulated proliferation of endothelial cells, modulating the proliferation of proliferation of proliferation of endothelial cells, inducing proliferation of cells, modulating glucose or FFA uptake, inducing proliferation and/or rectinal neurons or rod photoreceptor cells, inducing proliferation and/or redifferentiation of chondrocytes. In particular, these are useful for detecting or treating cardiac insufficiency disorders, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulatingenia,

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cutritis) in mammals. PRO polypeptides and their portions affect the arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides care useful in molecular biology including uses as hybridisation probes for conseil in chromosome and gene mapping, in the generation probes conseil in chromosome and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polypeptides, for generating transgenic confinals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals with genetic disorders as well as for recombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoreats purposes, as therapeutic agents, for screening compounds to identify the proteins are useful as molecular marker for protein electrophoreats to be seful for tissue typing. PRO antibodies are useful for tissue typing. PRO antibodies are useful for continual and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity continuals in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to maplify a PRO polymucleotide of the invention.
   hypoinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or
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Seguence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps . 0 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels Query Match 0.4 Best Local Similarity 93.8 Matches 15; Conservative

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1101 GCTGTCCTCAGGGGAG 1116 3 gererecacacacaca 18 à 셤

ADF73906 standard; DNA; 18 BP. RESULT 1946 ADF73906

26-FEB-2004 (first entry) ADF73906;

Human secreted/transmembrane protein, #44, PCR primer #2.

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; none disorder; cartilage disorder; cartilage disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological;

Homo sapiens.

osteopathic; antiarthritic; anorectic.

25-SEP-2003.

.8-NOV-2002; 2002US-00299976.

22-FEB-2000; 2000WO-US004414. 18-SEP-2000; 2000US-00665350.

(GETH) GENENTECH INC.

Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin LJ; Mather JP, Pan J, Paoni NP, Roy MA, Stewart TA, Tumas D;

Human secreted/transmembrane protein, #44, PCR primer #2.

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New PRO polypeptide useful for preparing a medicament for treating a condition that is responsive to the PRO polypeptide or anti-PRO antibody, e.g. inflammatory diseases, cancer or acquired immunodeficiency syndrome.
                                                                                                                                                                                                                                                                                           amplify a PRO polynucleotide of the invention.
                                           Example 36; SEQ ID NO 229; 473pp; English
Wood WI;
          WPI; 2004-031838/03.
Williams PM,
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27-0CT-1
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The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise and the nucleic acid encoding them. The PRO polypeptide for linking a bioactive molecule to a cell expressing a PRO polypeptide. For inking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide are useful for industry of a cell. PRO polypeptide are useful for modulating the biological activity of a cell. PRO polypeptide are useful for modulating the biological activity of conference of a cell expressing PRO pelypeptides. The PRO polypeptides are useful for stimulating the biological activity of a cell. PRO polypeptide on the properties are useful for stimulating the proliferation of polymorphy of neonatal provant factor (VEGP) stimulated proliferation of endochtalial cells, modulating the proliferation of retinulated proliferation of chondrocytes, chandring the survival or proliferation of chondrocytes, chandring the survival or proliferation and/or recills, modulating glucose or PRA uptake, inducing proliferation and/or recills, modulating glucose or PRA uptake, inducing proliferation and/or recills, modulating glucose or PRA uptake, inducing proliferation and/or recills, modulating or road photoreceptor cells, modulating modulating strangers or propression or propression of genes which beet with greates, hyperinalinaemia, cancerous cells, modulating glucose or PRA uptake, inducing proliferation and/or recills, modulating or bone or cartilage disorders (e.g. pports induced to the century or bone or cartilage disorders (e.g. pports induced to the century or bone or cartilage disorders (e.g. pports induced to the propression of genes which have useful in molecular biology including uses as hybridisation probes are useful in molecular marker for procein electrophores or purposes, as therapeutic agents of procein and plan with are useful in disorders are useful in disorders are useful as molecular marker for procein and propression in specific
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ADF73482 standard; DNA; 18

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(first entry)

26-FEB-2004 ADF73482;

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tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VBGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                   Human, PCR, primer, 88, PRO, secreted, transmembrane, therapeutic,
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970S-0059117P
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98US-0099803P
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                                                                                                                                   Homo sapiens.
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98US-0113296P.
99US-0143048P.
99US-0145698P.
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05-OCT-1999;
29-NOV-1999;
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Ferrara N;
1 ME, Goddard A;
Kljavin IJ;
Tumas D; Eaton DL, F I, Gerritsen M Hillan KJ, K Stewart TA, Ashkenazi A, Botstein D, Desnoyers L, Filvaroff E, Fong S, Gao W, Gerber H, Godowski BJ, Grimaldi JC, Gurney AL, H Mather JP, Pan J, Paoni NF, Roy MA, S Pan J, Pac Wood WI; William PM,

WPI; 2004-020549/02.

New secreted and transmembrane PRO polypeptides and nucleic acids, useful in gene therapy, in chromosome and gene mapping, as chromosome markers, in tissue typing, in identifying chromosomes, and for treating e.g. tumor or arthritis.

Example 36; SEQ ID NO 229; 478pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise action that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for inking a teast one biological activity of a cell. PRO polypeptides are useful concerned for linking a bloactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing a PRO polypeptides. The PRO polypeptides or polymenticals, diagnostics, biosensors or polymentical are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothelial growth factor (VEGF)-stimulated proliferation of chimulated T-lymphocytes, enhancing the survival or proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of cells, modulating glucose or FFA uptake, inducing proliferation and/or receiptor cells, inducing proliferation and/or receiptor cells, modulating closs in endothelial cells, cells, modulating proliferation and/or receiptor cells, inducing are useful for modulating chose or FFA uptake, inducing proliferation and/or receiptor cells, inducing are useful for modulating and control in particular, these are useful for modulating control in particular, these are useful for modulating control in particular.

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cc detecting or treating cardiac insufficiency disorders, wounds, cancerous tumours, retinal disorders or injuries (e.g. loss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arbitritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides or consistent biology including uses as hybridisation probes of or cDNA library to isolate the full-length PRO cDNA or to isolate other con DNA, for preparing PRO polypeptides, for generation of antisense RNA can DNA, for preparing PRO polypeptides, for generation of antisense RNA can and so therapeutically useful reagents, as probes and for the screening of therapeutically useful reagents, as probes and for the crecombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify the proteins are useful as molecular marker for protein electrophoresis purpose that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polymucleotides and proteins are useful for tissue typing. PRO antibodies are useful for emmunohistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic cassay of sample fluids. Anti-PRO purification of PRO from recombinant cassay of sample fluids. Anti-PRO purification of PRO from recombinant cassay of sample fluids sources. The polymucle and protein and protein
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nes 15; Conservative
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ADG92325 standard; DNA; 18 BP ADG92325; ADG92325

Human secreted/transmembrane protein, #44, PCR primer #2.

(first entry)

11-MAR-2004

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinaulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.

US2003027145-A1.

17-JUL-2001; 2001US-00907613 06-FEB-2003.

97US-0059113P. 97US-0059115P. 97US-0059117P. 97US-0059119P. 97US-0059121P. 17-SEP-1997; 7-SEP-1997 7-SEP-1997 7-SEP-1997 17-SEP-1997

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2000WO-US0084397. 2000WO-US0140439. 2000WO-US014042. 2000WO-US015264. 2000WO-US020710. 2000WO-US005004. 2000WO-US005841. 24-AUG-2000; 2000WO-US023328. 18-SEP-2000; 2000US-00665350. 24-FEB-2000; 2 02-MAR-2000; 2 20-MAR-2000; 3 30-MAR-2000; 2 22-MAY-2000; 2 05-JAN-2000; 11-FEB-2000; 22-FEB-2000; 28-JUL-2000;

(GETH) GENENTECH INC.

Ferrara N;
n ME, Goddard A;
Kljavin IJ;
Tumas D; Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NF, Roy MA, Stewart TA, Eaton DL, Desnoyers Botstein D, Ashkenazi A, Botstein I Filvaroff E, Fong S, G Godowski FU, Grimaldi G Mather Up, Pan J, Paor Williams PM, Wood WI,

WPI; 2004-118832/12.

New nucleic acid encoding a PRO polypeptide for use as hybridization probes, in chromosome and gene mapping, in generating antisense RNA and DNA, and in gene therapy for treating e.g. cancer, Parkinson's disease and wounds.

Example 36; SEQ ID NO 229; 471pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nurcleic acid enceding them. The polypeptides can be used to raise antichem uncleic acid enceding them. The polypeptides con be used to raise antichodise that specifically bind to the PRO polypeptide for inclining a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptides are useful for modulating the biological activity of a cell properties and sample and for inclining a bioactive molecule to a cell expressing PRO polypeptides are useful for modulating the biological activity of polypeptide antibodies are useful for modulating the polypeptides or polymuclectides are useful for stimulating hypertrophy of neonatal collinearing of a cell expressing PRO polypeptides. The PRO polypeptides or polymuclectides are useful for stimulating hypertrophy of neonatal carriers or stimulated properties can be accepted to the polymuclectides are useful for stimulating hypertrophy of neonatal carriers or red plantaneous or rod phorocreceptor cells, modulating policose or FRA uprace, inducing proliferation of chondrocryes. Handle properties are useful for cells, modulating plucose or FRA uprace, inducing proliferation of chondrocryes. Inducing proliferation of chondrocryes in particular, these are useful for cells, modulating plucose or FRA uprace, inducing proliferation of chondrocryes or injuries (e.g. loss of sight due to carriers or retinate properties or injuries (e.g. loss of sight due to continuous or to appropriate or properties and their portions affect the arbitration of genes which have a role in cell for cupy library to isolate the fell. In molecular bloopy including uses as hybridisation probes are useful in molecular bloopy including uses as hybridisation probes or control of solate the fell of the properties of the protein and to a role of the process, as therapeutically useful respents of solates are useful for the properties of individuals with generic antipulate or the properties of individuals with generic an

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Botstein D, Desnoyers L, Eaton DL, Ferrara N; Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; an J, Paoni NF, Roy MA, Stewart TA, Tumas D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid encoding a polypeptide useful for various applications e.g. hybridization probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JAN-2000; 2000WO-US000219.
11-FEB-2000; 2000WO-US003565.
22-FEB-2000; 2000WO-US004414.
24-FEB-2000; 2000WO-US005004.
02-MAR-2000; 2000WO-US005841.
20-MAR-2000; 2000WO-US005841.
       97US-0063738P.
97US-0064215P.
97US-0064103P.
97US-0064103P.
97US-0065186P.
97US-0065186P.
97US-0065186P.
97US-0066120P.
97US-0066453P.
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97US-0109304P.
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98US-010480P.
98US-010480P.
99US-0145698P.
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99WO-US028313.
99WO-US028301.
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99WO-US028565.
99WO-US030095.
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99WO-US021090.
99WO-US021547.
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24-AUG-2000; 2000WO-US023328.
18-SEP-2000; 2000US-00665350.
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Filvaroff E, Fon
Godowski PJ, Gri
Mather JP, Pan J
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                                      31-OCT-1997;
03-NOV-1997;
07-NOV-1997;
                                                                     12-NOV-1997;
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02-DEC-19
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ö
                                                                                                                                                                                                                                    tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; cod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; oardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritis; anorectic.
                                                  Gaps
                                                                                                                                                                                                                                Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic;
                                                  ;
0
                           Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                           Human secreted/transmembrane protein, #44, PCR primer #2.
         Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                     1101 GCTGTCCTCAGGGGAG 1116
                                                                                                                                                                                                                                                                                                                                                                                                                             9705-0059113P
9705-0059113P
9705-0059113P
9705-0059121P
9705-0059122P
9705-0059122P
9705-0059124P
9705-0062285P
9705-0062285P
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9705-0062285P
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9705-0063542P
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9705-0063542P
9705-0063542P
9705-0063542P
9705-0063542P
                                                                                                                                             ADG92752 standard; DNA; 18 BP.
                                                                                  GCTGTCCACAGGGGAG 18
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                                                                                                                                                                                     11-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                  US2003027146-A1
                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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17.58P-1997
17.58P-1997
18.58P-1997
18.58P-1997
17.0CT-1997
17.0CT-1997
24.0CT-1997
24.0CT-1997
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24-0CT-1997;
24-0CT-1997;
27-0CT-1997;
27-0CT-1997;
28-0CT-1997;
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29-OCT-1997;
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28-OCT-1997
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28-OCT-1997
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98WO-US019437
98US-0104080P
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                                                                           Homo sapiens.
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27-0CT-
 The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucled acid encoding them. The polypeptides contribodies that specifically bind to the PRO polypeptide for linking a bioactive molecule to a cell expressing a PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO polypeptides are useful contribodies are useful for modulating the biological activity of a cell. PRO polypeptides or a cell expressing a PRO polypeptides or bolypeptides are useful for modulating the biological activity of a cell expressing a PRO polypeptides. The PRO polypeptides or polypeptides are useful for modulating the biological activity of a cell expressing as pharmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating the proliferation of endothelial cells, modulating the proliferation of proliferation of condothelial cells, modulating the proliferation of stimulated T-lymphocytes, enhancing the survival or proliferation of condothelial cells, modulating proliferation of conductoreceptor cells, inducing proliferation of conductoryes. In particular, these are useful for cetinal neurons or rod photoreceptor cells, inducing proliferation of conductoryes. In particular, these are useful for cetinal induces or retained isorders or injuries (e.g. loso feight due to detecting or treating carders or injuries (e.g. loso feight due to cetinities pigmentosum), obesity, disorders (e.g. sports sinjuries or cetinities pigmentosum), obesity, disorders (e.g. sports sinjuries or ensetul in molecular biology including uses as hybridiation of genes which have a role in cell death. The polymucleotides are useful in molecular biology including uses as hybridiation of antisense RNA arrivals or thormosome and gene mapping, in the generating transgenic combinantly expression of genes which have a role in cell death. The polymucleotides are useful animals with genetic disorders as well animals or knockout animals with genetic analysis of individuals with genetic anophysis of individua
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; PCR; primer, ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucoses; PFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.4; DB 1; Length 18; 13.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
Example 36; SEQ ID NO 229; 474pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1101 GCTGTCCTCAGGGGAG 1116
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Matches 15; Conservative
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DT 25-MAR-2
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Human se
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98US-0113296P.
99US-0143048P.
99US-0145698P.
99US-0146222P.
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24-FEB-2000; 2000WO-US005004.
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                                                                                                                                                                                Wood WI;
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                                                                                                                                                      (GETH ) GENENTECH INC.
                                                                                                                                                                                            WPI; 2004-081703/08.
                                                                           16-DEC-1999;
20-DEC-1999;
20-DEC-1999;
05-JAN-2000;
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13-SEP-1999;
15-SEP-1999;
15-SEP-1999;
                                               05-OCT-1999;
29-NOV-1999;
30-NOV-1999;
                  26-JUL-1999
                                                                                                                                                                                                               typing
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Ferrara N; n ME, Goddard A; Kljavin IJ; Tumas D; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Flyaroff E, Fong S, Gao W, Gerber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA,

New PRO nucleic acid, useful for manufacturing a medicament for diagnosing or treating tumor, for chromosome mapping or for tissue

Example 36; SEQ ID NO 229; 126pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or useful as pharmaceuticals, diagnostics, biosensors or polymicleotides are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothelial growth factor (VGEP)-stimulated complexation of endothelial cells, modulating the proliferation of cells, modulating the proliferation of cells, modulating the proliferation of cells, modulating the survival or proliferation of cells, modulating the survival or proliferation of cells, modulating cardiac insufficiency disorders, wounds, cancerous cells, modulating cardiac insufficiency disorders, wounds, cancerous cummuns, retinal disorders or injuries (e.g. sports injuries or retinitis pigmentosum), obesity, diabetes, hyperinaulinaemia, creating cardiac insufficiency disorders, wounds, carcindes arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a roole in cell death. The polymuclectides are useful in melecular biology including uses as hybridisation probes for connactive the full-length PRO cDNA or to isolate other

New secreted and transmembrane PRO polypeptides and nucleic acids, useful

WPI; 2004-081748/08.

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and DNA, for preparing PRO polypeptides, for generating transgenic animals or knockout animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals with genetic disorders as well as for recombinantly expressing the protein and for chromosome identification.

The proteins are useful as molecular marker for protein electrophoresis or purposes, as therapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polynucleotides and proteins are useful for tissue typing. PRO antibodies are useful for tissue typing. PRO antibodies are useful for munohistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity comprehence or province therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
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in chromosome and gene mapping, in the generation of antisense RNA
                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; typing; immunohistochemical staining; gene therapy;
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                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein, #44, PCR primer #2.
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                                                                                                                                                                                                                                                                                                Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADH07396 standard; DNA; 18 BP
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18-SEP-2000; 2000US-00665350.
12-JUL-2001; 2001US-00905125.
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99WO-US028313.
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                                                                                                                                                                                                                                                                                                                                            93.8%;
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Wood WI;
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GURNEY A L.
MATHER J P.
WILLIAMS P M.
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Williams PM,
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15-OCT-1997;
17-OCT-1997;
24-OCT-1997;
25-OCT-1997;
26-OCT-1997;
27-OCT-1997;
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   The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bloactive molecule to a cell expressing a PRO polypeptide are useful cor modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide are useful as pharmaceuticals, diagnostics, blosensors or bloreactors. The PRO sequences can be used in gene and protein therapy. The PRO polypeptide, the agonist or antagonist or the anti-PRO antibody. The nucleic acids encoding PRO polypeptides are used as hybridisation probes for gene mapping, can tangenis are used as hybridisation probes for gene mapping. The PRO polypeptides are used as hybridisation probes for gene mapping. The PRO polypeptides are used as hybridisation or for tissue typing. The PRO polypeptides are also useful in gene therapy, may be employed as molecular weight markers for protein electrophoresis or as therapeutic and an are also useful in gene therapy, and the preparents and an are also useful in gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                  agents. Anti-PRO antibodies are useful in diagnostic assays or for the affinity purification of PRO for recombinant cell culture or natural sources. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.
 gene therapy, as molecular weight markers for protein electrophoresis, hybridization probes or as therapeutic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; nenonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmencosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartiage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; ive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                          Example 36; SEQ ID NO 229; 466pp; English.
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97US-0059115P.
97US-0059117P.
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Best Local Similarity 93.8
Matches 15, Conservative
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17-SEP-1997;
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PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polynucleotide of the invention.

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invention discloses isolated PRO secreted/transmembrane polypeptides
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Kljavin IJ;
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W, Gerber H, Gerritsen I
Gurney AL, Hillan KJ, 1
NF, Roy MA, Stewart TA,
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                                                                                                                                                            2000WO-US008439.
2000WO-US014042.
2000WO-US015264.
2000WO-US020710.
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2000WO-US007377.
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99WO-US030911
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Mather JP, Pan J, Pac
Williams PM, Wood WI;
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Filvaroff E,
                                                         11-FEB-2000;
22-FEB-2000;
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continuon discloses isolated PNO Secreted/Crabinembrane Polypoptides and the murchact acid encoding them. The polypoptides can be used to raise antibodies that specifically bind to the PRO portpeptide for linking a bioactive molecule to a cell expressing a PRO polypeptide. The RD polypeptides are useful expressing a PRO polypeptide. The PRO polypeptides are useful expressing a PRO polypeptide. The PRO polypeptides are useful expressing a PRO polypeptide. The PRO polypeptides or bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptides are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymeticotides are useful for stimulating hypertrophy of monatal bioractors. These are useful for stimulating hypertrophy of monatal cheart, inhibiting vascular endothelial growth factor (VBGP-stimulated profileration of endothelial cells, modulating the proliferation of control profileration of control proceeptor cells inducing cries in endothelial cells. modulating the proliferation of control cells modulating proliferation of control cells inducing refers inducing proliferation of control cells. modulating the survival or proliferation of celliferentiation of chondrocytes. In particular, these are useful for detecting or treating cardiac insufficiently there are useful for cetinal induced or repeating cardiac insufficiently describes (e.g. polymental) celliferentiation of genes which have a role in cell deach. The polymented cardiac insufficient or celliferent portions affect the articular biology including uses as hybridisation probes for control genes which have a role in cell death. The polymented and cellifered cardiac insurance and gene mapping, in the generating transgenic and mains or knockout annials which have a role in cell death. The generating of therapeutically useful reagents, for generating transgenic combinantly expressing the protein and control and properses as therapeutically useful reagents, or propersed to manipais of indiv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New PRO polypeptides and nucleic acid molecules, useful in gene therapy, or preparing a medicament for treating a condition that is responsive to the PRO polypeptide or anti-PRO antibody, e.g. inflammatory diseases, cancer or AIDS.
                                                                                                            Gaps
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                                                                              Score 14.4; DB 1; Length 18;
Pred. No. 1.7e+03;
0; Mismatches 1; Indels
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                                                    Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                      0.4%; Scor-
93.8%; Pred
0; V
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16-SEP-1998; 98WO-US019330.
30-NOV-1999; 99WO-US028313.
22-FEB-2000; 2000WO-US04414.
18-SEP-2000; 2000US-00665350.
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Wood WI;
                                                                                              Best Local Similarity 93.8 Matches 15; Conservative
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MATHER J P.
WILLIAMS P M.
WOOD W I.
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Williams PM,
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                        24-0CT-1997;
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    polymucleotides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. The PRO sequences can be used in gene and protein therapy. The PRO polypeptide, the agonist or antagonist or the anti-PRO antibody can be used in the preparation of a medicament for the treatment of a condition which is responsive to the PRO polypeptide, the agonist or antagonist or the anti-PRO antibody. The nucleic acids encoding PRO polypeptides are used as hybridisation probes for gene mapping, generating transgenic animals useful in the development and screening of useful reagents, in chromosome identification or for tissue typing. The PRO polypeptides are also useful in gene therapy, may be employed as molecular weight markers for protein electrophoresis or as therapeutic agents. Anti-PRO antibodies are useful in diagnostic assays or for the sequence presented is a PCR primer which was used to amplify a PRO polymucleotide of the invention.
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                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                   0.4%; Score 14.4; DB 1; Length 18; 13.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                                                                              Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                   1101 GCTGTCCTCAGGGGAG 1116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9705-0059115P
9705-0059117P
9705-0059119P
9705-0059121P
9705-0059184P
9705-0059264P
9705-0059268P
9705-0062818P
9705-0062818P
9705-0062818P
9705-0062816P
9705-0062816P
9705-0062816P
                                                                                                                                                                                                                                                                                                                                                     ADI18711 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                           GCTGTCCACAGGGAG 18
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                                                                                                                                                                                                                                 93.8%;
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                                                                                                                                                                                                                               Local Similarity 93.8
es 15, Conservative
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17-SEP-1997
17-SEP-1997
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17-SEP-1997
18-SEP-1997
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11-OCT-1997
17-OCT-1997
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24-OCT-1997;
24-OCT-1997;
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Matches
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18-SEP-2000; 2000US-00665350

Human secreted/transmembrane protein, #44, PCR primer #2.

(first entry)

22-APR-2004

ADI65431;

Ferrara N; ME, Goddard A; Kljavin IJ;

ADI65431 standard; DNA; 18 BP

RESULT 1955

ADI6543

Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, I ElVaroff E, Fong S, Gao W, Gerber H, Gerriteen P Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, I Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, (GETH) GENENTECH INC. Godowski PJ, Grimaldi Mather JP, Pan J, Pa Williams PM, Wood WI;

WPI; 2004-020479/02,

Sixty two isolated nucleic acids encoding a PRO polypeptide, e.g. PRO245 or PRO1868, useful for treating psoriasis and epithelial cancers such as lung squamous cell carcinoma.

Example 36; SEQ ID NO 229; 426pp; English.

The invention is decided them. The polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a to loactive molecule to a cell expressing a PRO polypeptide, for linking a to loactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide in the propertide at least one biological activity of a cell. PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of polypeptide antibodies are useful for modulating the biological activity of polypeptide antibodies are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endethelial growth factor (VEGP)-stimulated contributing vascular endethelial growth factor (VEGP)-stimulated contributing vascular endethelial growth factor (VEGP)-stimulated contributing or or polymerceeptor cells, inducing c-fos in endothelial cells, modulating proliferation of chondrocytes, enhancing the survival or proliferation of cellman activity and proliferation of chondrocytes. In particular, these are useful for retinal neurons or rob photoreceptor cells, inducing proliferation and/or reciting and chordrocytes. In particular, these are useful for cells, modulating glucose or PFA uptake, inducing proliferation and/or reciting in modulating glucose or PFA uptake, inducing proliferation of chondrocytes. In particular, these are useful for centuring of chondrocytes. In particular, the polymerical disponses or injuries (e.g. loss of sight due to retinities) in ammals, or bone or cartilage disorders (e.g. sports injuries or retinities) in ammals, and polymerical death. The polymerical control of genes which have a role in cell death. The polymerical control of an individuals with genetic disorders as well as for conditional, which are useful in the development and screening of therapeutically useful reagents, as probes as therapeutic analysis of individuals with genetic and part of process as therapeutic agen antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to The invention discloses isolated PRO secreted/transmembrane polypeptides defective gene. The sequence presented is a PC amplify a PRO polynucleotide of the invention.

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

; 0 0.4%; Score 14.4; DB 1; Length 18; 33.8%; Pred. No. 1.7e+03; 1; Indels 0; Mismatches 93.8%; Best Local Similarity 93.8 Matches 15; Conservative Query Match

1101 GCTGTCCTCAGGGGAG 1116

상· 원

3 gererecacadedeag 18

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Gaps

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; enconatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diábece; hyperinaulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic. 9705-00631219-9705-0063128P-9705-0063327P-9705-0063329P-9705-0063541P-9705-0063544P-9705-006354P-9705-006354P-9705-0063564P-970S-0059119P. 970S-0059121P. 970S-0059184P. 970S-0059263P. 970S-0059266P. 97US-0062816P. 97US-0063045P. 97US-0063120P. 970S-0063435P. 970S-0063704P. 970S-0063732P. 970S-0063734P. 970S-0063735P. 97US-0063870P. 97US-0064103P. 97US-0064248P. 97US-0064809P. 97US-0063486P. 97US-0062814P. 11-JUL-2001; 2001US-00903603 97US-0059113P 97US-0063738P 97US-0064215P 97US-0066120P 97US-0066364P US2003148419-A1. Homo sapiens. 07-AUG-2003. 21-NOV-1997; 997 29-0CT-1 29-0CT-1 29-OCT-1 27-OCT-: 28-0CT

diagnostics, biosensors or

vivlemore401-10.rng

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97US-0066466P
                                  97US-0069425P
                             97US-0066840P
                                                    98WO-US018824
                                                                           98US-0100858P
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24-FEB-2000;
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24-NOV-1997;
24-NOV-1997;
24-NOV-1997;
24-NOV-1997;
25-NOV-1997;
                                       04-JUN-1998;
10-SEP-1998;
10-SEP-1998;
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15-OCT-1999;
29-NOV-1999;
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                                                         14-SEP-1998
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13-SEP-1999
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(GETH) GENENTECH INC.

ME, Goddard A; Kljavin IJ; Tumas D; Ferrara N; Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Filvaroff E, Pong S, Gao W, Gerber H, Gerritsen Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Wood WI: Williams PM,

WPI; 2004-020444/02.

New isolated secreted and transmembrane PRO nucleic acids and polypeptides, useful for preventing, diagnosing and treating disorders associated with their aberrant expression and activity.

Example 36; SEQ ID NO 229; 476pp; English.

and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polymentides can he used to raise

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C polymuclacotides are useful as pnarmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating hypertrophy of neonatal carbinating avacular endothelial growth factor (WEGP)-stimulated proliferation of endothelial cells, modulating the proliferation of endothelial cells, modulating the proliferation of the stimulated T-lymphocytes, enhancing the survival or proliferation of control processors in ducing proliferation of retinal neurons or rod photoreceptor cells, inducing proliferation of cells, modulating glucose or FRA uptake, inducing proliferation and/or recombination of chondrocytes. In particular, these are useful for detecting or treating cardiac insufficiency disorders wounds, cancerous cutmours, retinal disorders or injuries (e.g. loss of sight due to detecting or treating cardiac insufficiency disorders wounds, cancerous cutmours, retinal disorders or injuries (e.g. ports injuries or arthitis) in mammals. PRO polyeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides are useful in molecular blooky including uses as hybridisation probes care useful in molecular and gene mapping, in the generation probes convince of the proposition and gene mapping, in the generation of antisense RNA and DNA, for preparing PRO polyeptides, for generation transgenic animals or knockout animals which are useful in the development and corrections of therspeutically useful reagents as probes and for the combinantly expressing the protein and for chromosome identification. The proteins are useful in molecular marker for protein electrophoresis purposes, as therapeutic adents, for screening compounds to identify the protein are useful for the protein and for chromosome identification. The proteins are useful in diagnostic assays for protein electing its correlating or purposes, as therapeutic adents, for screening compounds to identify the protein and protein and for the protein and protein and protein and protein and protein and protein and pro
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Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps .; 0 Match 0.4%; Score 14.4; DB 1; Length 18; Local Similarity 93.8%; Pred. No. 1.7e+03; les 15; Conservative 0; Mismatches 1; Indels Query Match Matches

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RESULT 1956 ADI82699

ADI82699 standard; DNA; 18 AD182699;

Vector pEGFP(His) DNA fragment #4.

(first entry)

22-APR-2004

lysis; cytoskeleton; EF-Tu; bacterial elongation factor; fibril assembly; protofilament; cell wall rupture; ds.

Synthetic.

WO2004005506-A2.

15-JAN-2004.

02-JUL-2003; 2003WO-EP007068.

02-JUL-2002; 2002DE-01029645.

(NOVO-) NOVOLOGIX GMBH

Schwienhorst A; Mayer F,

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9705-0059184P
9705-005926B
9705-005926B
9705-0062187P
9705-0062187P
9705-0062816P
9705-0062816P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063121P
9705-0063131P
9705-006411P
9705-0066453P
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98US-0109304P.
98WO-US025108.
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99US-0143048P.
99US-0145698P.
99US-0146222P.
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99WO-US030095
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05-OCT-1999
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24-0CT-1997
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21-0CT-19
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29-0CT-1
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31-0CT-1
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This invention describes a novel method for lysing cells using a substance that binds to components of the cytoskeleton, especially to EF-
c substance that binds to components of the cytoskeleton, especially to EF-
c Tu (bacterial elongation factor), destabilising it. The substance binds
to domains 2 and/or 3 (respectively, amino acids 205-298 and 299-349) of
ET-tu, especially to regions 218-24, 317-328 and/or 343-354. Preferred
c EF-tu, especially to regions 218-24, 317-328 and/or 343-354. Preferred
c Inhibiting self-assembly of EF-Tu fibrils. The polymucleotide encoding
the cytoskeleton-binding substance is in a construct that also includes
the polymucleotides that induce its synthesis. The products of the
c invention are used to lyse bacteria, specifically where these are being
c used to prepare heterologous or homologous intracellular compounds,
c used to prepare heterologous or homologous intracellular compounds,
c especially proteins for biotechnological or medical applications.
C Escherichia coli cells, containing a gene that encoded the factor EF-Tu,
c were transformed with a plasmid that expressed domain 3 of EF-Tu. This
domain competes with complete EF-Tu in the formation of protofilaments
c and where incorporated it causes a strand break (it contains no domain 2,
c which is essential for chain formation). Defective protofilaments can not
c grow, so the cytoskeletal network is weakened and collapses. This removes
the support for the cytoplasmic membrane, resulting in rupture of the
c lis used in the method described in the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; cod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                           Inducing cell lysis with compound that binds to a cytoskeletal component, useful for recovering homologous or heterologous proteins from bacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;
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                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

0.4%; Score 14.4; DB 1; Length 18;
Best Local Similarity 93.8%; Pred. No. 1.7e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein, #44, PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                     Disclosure; Fig 3; 36pp; German.
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97US-0059115P.
97US-0059117P.
97US-0059119P.
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WPI; 2004-091366/09.
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17-SEP-1997;
17-SEP-1997;
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Gaps

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1; Indels

Score 14.4; DB 1; Pred. No. 1.7e+03; 0; Mismatches 1;

0.4%;

3 GCTGTCCACAGGGAG 18

Length 18;

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New PRO nucleic acid, useful for producing a PRO polypeptide, manufacturing a medicament for diagnosing or treating tumor, or for
                                                WPI; 2004-008942/01.
                                                         tissue typing
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Human secreted/transmembrane protein, #44, PCR primer #2.
                                   Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
amplify a PRO polynuclectide of the invention.
                                                                                                                                                  1101 GCTGTCCTCAGGGGAG 1116
                                                                                                                                                                                                                                                                                   ADH97490 standard; DNA; 18 BP
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                                                                        Query Match 0.4
Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                         ADH97490;
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Kljavin IJ;
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Hillan KJ, K
Stewart TA,
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W, Gerber H,
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                                                                      22-FEB-2000; 2000WO-US00414.
24-FEB-2000; 2000WO-US005004.
02-MAR-2000; 2000WO-US005841.
                                                                                                                                                                                     02-JUN-2000; 2000WO-US015264.
28-JUL-2000; 2000WO-US023128.
24-JUC-2000; 2000WO-US023328.
18-SEP-2000; 2000US-00665350.
                                                                                                                               20-MAR-2000; 2000WO-US00777
30-MAR-2000; 2000WO-US008439.
22-MAY-2000; 2000WO-US014042
   99WO-US030911
                   99WO-US030999
2000WO-US000219
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Godowski PJ, Grimaror
Mather JP, Pan J, Pac
                                                                                                                                                                                                                                                                                     (GETH ) GENENTECH INC.
20-DEC-1999;
20-DEC-1999;
05-JAN-2000;
11-FEB-2000;
                                                                                                                                                                                                                                                                                                                         Ashkenazi A,
Filvaroff E,
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Example 36; SEQ ID NO 229; 474pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise articodises that specifically bind to the PRO polypeptide. To linking a to bioactive molecule to a cell expressing a PRO polypeptides are useful controlled expressing PRO polypeptides. The PRO polypeptides are useful controlled expressing PRO polypeptides. The PRO polypeptide articolled expressing PRO polypeptides. The PRO polypeptide articolled expressing PRO polypeptides. The PRO polypeptides or colled expressing PRO polypeptides or colled expressing PRO polypeptides. The PRO polypeptides or colled expressing PRO polypeptides. The PRO polypeptides or useful for stimulating hypertrophy of neonatal colled expressing PRO polypeptides. The PRO polypeptides or colled proliferation of endothalial cells, modulating the proliferation of endothalial cells, modulating the proliferation of endothalial cells, modulating proliferation of endothalial cells, modulating proliferation of endothalial cells, inducing proliferation of colled procreceptor cells, inducing proliferation of endothalial cells, inducing proliferation of colled colled colled proliferation of colled procreceptor cells, inducing proliferation of colled colled colled proliferation of colled co purposes, as therapeutic agents, for screening compounds to identify those that mimic the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polymoleotides and proteins are useful for tissue typing. PRO antibodies are useful for immunohistochemical staining and/or assay of sample fluids. Anti-PRO antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to

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tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; cod photoreceptor cell; cfos; glucose; fish, chondrocyte; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperineulinaemia; hypoineulinaemia; hypoineulinaemia; one disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
                                                                                                                                                                                                       Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;
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97US-0059117P.
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9703-0063564P.
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97US-0063738P.
97US-0064215P.
97US-0063870P.
97US-0064248P.
97US-0064248P.
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Goddard A; ME, Goddar Kljavin IJ; Ferrara N; Tumas D; Gerritsen M Hillan KJ, K Stewart TA, Eaton DL, Gerber H, Fong S, Gao W, Gerber H Grimaldi JC, Gurney AL, an J, Paoni NF, Roy MA, Botstein D, Desnoyers L, Wood WI; Pan J, Ashkenazi A, Filvaroff E, Godowski PJ, Mather JP, P? Williams PM,

WPI; 2004-032142/03.

New nucleic acid encoding a PRO polypeptide, useful for producing a recombinant PRO polypeptide and for treating tumors by gene therapy.

Example 36; SEQ ID NO 229; 471pp; English.

C The invention discloses isolated RNO secreted/transmembrash polypeptides can be used to raise and the nucleic acid encoding them. The polypeptides can be used to raise can the invention discloses isolated RNO secreted/transmembrash for inhiding a bloative molecule to a cell expressing a PRO polypeptide for inhiding a condition other PRO polypeptides in a sample and for linking a bloative molecule to a cell expressing a PRO polypeptide. The PRO polypeptides are useful as pharmaceuticals, diagnostics, biosensors or polypeptides are useful in paramaceuticals, diagnostics, biosensors or polypeptides are useful as pharmaceuticals, diagnostics, biosensors or polypeptides are useful as pharmaceuticals, diagnostics, biosensors or polymelocides are useful appraismenticals, diagnostics, biosensors or proliferation of endothelial cells, modulating the proliferation of endothelial cells, modulating the proliferation of endothelial cells, inducing proliferation of endothelial cells, inducing proliferation and/or cells, modulating diacose or FRA uptaske, inducing proliferation and/or cells, modulating or treating cardiac insufficiency disorders, wounds, cancerous cells, modulating or bone or cartiloge disorders, hyperinsulation of chondrocytes, inquiries (e.g. loss of sight due to creintis pigmentosum) obesity, diaberes, hyperinsulinaemia, hypoinaulinaemia, or bone or cartiloge disorders (e.g. loss of sight due to creintis pigmentosum) obesity, diaberes, hyperinsulinaemia, chopy including use as hybridisation probes or crimitis pigmentosum) obesity, diaberes, hyperinsulinaemia, chopy including use as hybridisation probes or crimitis pigmentosum) obesity, diaberes, hyperinsulinaemia, and of certaperating PRO polypeptides, for generating transgenic and mals or knockout animals which are useful in the generation of antisense RNA are suffered in molecular bology including use as hybridisation process, as theraputic agents, for screening of theraperating PRO polypeptides, for generating transgenic and public cannysis of individ The invention discloses isolated PRO secreted/transmembrane polypeptides defective gene. The sequence presented is a posemplify a PRO polynucleotide of the invention.

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match 0.4%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 1.7e+03; Matches 15; Conservative 0; Mismatches 1; Indels

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1101 GCTGTCCTCAGGGGAG 1116 GCTGTCCACAGGGGAG 18 ò

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ADI65858 standard; DNA; 18 BP. ADI65858; RESULT 1959 AD165858

22-APR-2004 (first entry)

Human secreted/transmembrane protein, #44, PCR primer #2.

Human; PCR; primer, ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VSGF; proliferation; endothelial cell; stimulated Tlymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte;

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cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic;
                                                                                                                 97US-0059113P.
97US-0059113P.
97US-0059117P.
97US-0059121P.
97US-0059121P.
97US-0059124P.
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98US-0088026P.
98US-0099803P.
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                                                   Homo sapiens.
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98US-0113296P.
99US-0143048P.
99US-0145698P.
99US-0146222P.
99WO-US020594.
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99WO-US028301.
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2000WO-US023328
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                                                                                    22-FEB-2000;
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29-NOV-1
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Ferrara N; ME, Goddard A; Kljavin IJ; Tumas D; Eaton DL, E , Gerritsen N Hillan KJ, F Stewart TA, Desnoyers L, W, Gerber H, ldi JC, Gurney AL, Paoni NF, Roy MA, Fong S, Gao W, Grimaldi JC, G Ashkenazi A, Botstein D, Filvaroff E, Fong S, Gao Godowski PJ, Grimaldi JC, Mather JP, Pan J, Paoni N Williams PM, Wood WI;

WPI; 2004-020441/02.

Isolated secreted and transmembrane PRO nucleic acids and the proteins they encode, e.g. PRO245, PRO269 and PRO1868, useful for preventing, diagnosing and treating e.g. disorders relating to blood coagulation.

Example 36; SEQ ID NO 229; 478pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bloactive molecule to a cell expressing a PRO protein and for modulating a conference on other PRO polypeptides in a sample and for linking a conference of the properties of the properties are useful conference of a cell expressing PRO polypeptides or the PRO polypeptides or coll expressing PRO polypeptides or collection of collection of collection of collection and collection of collection collection collection of collection of collection collection collection collection of collection col

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and DNA, for preparing PRO polypeptides, for generation of antisense RNA animals which are useful in the development and screening of therapeutically useful reagents, as probes and for the genetic analysis of individuals with genetic disorders as well as for recombinantly expressing the protein and for chromosome identification. The proteins are useful as molecular marker for protein electrophoresis purposes, as therapeutic agents, for screening compounds to identify those that miniar the PRO polypeptide (agonists) or prevent the effect of the PRO polypeptide (antagonists). The polymucleotides and proteins are useful for tissue typing. The polymucleotides and proteins are useful for tissue typing and/or assay of sample fluids. Anti-PRO antibodies are useful for antibodies are useful for captession in specific cells, tissues or serum and for affinity expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a defective gene. The sequence presented is a PCR primer which was used to amplify a PRO polymucleotide of the invention.
cDNA library to isolate the full-length PRO cDNA or to isolate other
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0; Mismatches 1101 GCTGTCCTCAGGGAG 1116 93.8%; 15; Conservative Sest Local Similarity Query Match fatches

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Human secreted/transmembrane protein, #44, PCR primer #2. ADH60601 standard; DNA; 18 BP. (first entry) 22-APR-2004 ADH60601;

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartliage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.

Homo sapiens.

28-APR-2003; 2003US-00425447 97US-0063128P 24-OCT-1997;

GODOWSKI P J. GURNEY A L. MATHER J P. WILLIAMS P M. GODDARD A. (DESN/) 1 (GODD/) (GODD/) (GURN/) (MATH/) 1 (WILL/) 1

Gaps .; 0 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; 1; Indels Seguence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

3 GCTGTCCACAGGGAG 18 ઠે g

RESULT 1960
ADH60601
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AC ADH60601
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DT 22-APR-2
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Human 8
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Human 1, 18
XW
Human 2, 18
XW
Human 3, 18
XW
Cardiac

US2004023331-A1. 05-FEB-2004.

16-SEP-1998; 98WO-USO19330. 30-NOV-1999; 99WO-USO28313. 22-FEB-2000; 2000WO-USO04414. 18-SEP-2000; 2000US-00665350. 17-JUL-2001; 2001US-00907794. DESNOYERS L.

(WOOD/) WOOD W I.

Gurney AL, Mather JP; Godowski PJ, Goddard A, Wood WI; Desnoyers L, Williams PM,

WPI; 2004-142655/14.

New secreted and transmembrane nucleic acids and polypeptides, designated as PRO, useful for treating inflammation, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, or

Disclosure; SEQ ID NO 229; 428pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nutcleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide. For linking a cariobacise that specifically bind to the PRO polypeptide. For linking a taleast one blooks are useful for a cell sepressing a PRO polypeptide The PRO polypeptide antibodies are useful for modulating the biological activity of a cell. FRO polypeptide antibodies are useful for modulating the biological activity of a cell. FRO polypeptide on the PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptide in the PRO polypeptide antibodies are useful for modulating properties, biosensor or polymuclectides are useful for stimulating hypertrophy of meaners or polymuclectides are useful for stimulating hypertrophy of meaners or polymuclectides are useful for stimulating proliferation of chondrocytes, inducing proliferation of chondrocytes, inducing proliferation of chondrocytes, inducing proliferation of chondrocytes. In particular, these are useful for calification of chondrocytes. In particular, these are useful for calification of chondrocytes or injuries (e.g. loss of sight due to creintain or dense which have a rote in media. Propring the creinties promines are useful in molecular hology including uses as hybridisation probes or prepagation of genes which have a rote in call death. The polymuclecties are useful in molecular hology including uses as hybridisation probes or on DNA, in chromosome and gene mapping, in the generating rangement and control of the properties are useful in molecular hology including uses as probes and for the generating repetition of chondrocyte animals or knockout animals which are useful in the development and gene mapping, in the generating cranged canneral or recombinantly expressing the protein and for chromosome and gene mapping or creaming one protein are useful as molecular marker for protein electrophores; or

Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

Gaps . 0 0.4%; Score 14.4; DB 1; Length 18; 3.8%; Pred. No. 1.7e+03; ve 0; Mismatches 1; Indels 93.8%; Query Match 0.4 Best Local Similarity 93.8 Matches 15, Conservative

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1101 GCTGTCCTCAGGGGAG 1116

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BP. ADJ99658 standard; DNA; 18 RESULT 1961 ADJ99658

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tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; hypoinsulinaemia; oardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritis; anorectic.
                                      Human; PCR; primer; 88; PRO; secreted; transmembrane; therapeutic;
                          Human secreted/transmembrane protein, #44, PCR primer #2
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99US-0143048P.
99US-0145698P.
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11-FEB-2000;
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24-FEB-2000;
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Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N; Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A; Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ; Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D; Williams PM, Wood WI;

WPI; 2004-032054/03.

Isolated nucleic acid for making vector for host cell, comprises specified sequence identity to nucleotide sequence that encodes polypeptide having amino acid sequence.

Example 36; SEQ ID NO 229; 470pp; English.

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO polypeptide, for linking at least one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptide. The PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polypeptides are useful as pharmaceuticals, diagnostics, biosensors or bioreactors. These are useful for stimulating hypertrophy of neonatal heart, inhibiting vascular endothelial growth factor (VEGF) estimulated proliferation of endothelial cells, modulating the proliferation of

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       cells, modulated T-lymphocytes, enhancing the survival or proliferation of cells, inducing c-fos in endothelial calls, modulating glucose or FRA uptake, inducing c-fos in endothelial calls, modulating glucose or FRA uptake, inducing proliferation and/or reclifferentiation of chondrocytes. In particular, these are useful for detecting or treating cardiac insufficiency disorders, wounds, cancerous turnurs, retinal disorders or injuries (e.g. 10ss of sight due to retinitis pigmentosum), obesity, diabetes, hyperinsulinaemia, or bone or cartilage disorders (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides care useful in molecular biology including uses as hybridisation probes of for companient of genes which have a role in cell death. The polymucleotides companient of companient and DNA, for preparing PRO polypeptides, for generation of antisense RNA and DNA, for preparing PRO polypeptides, for generation of antisense RNA and DNA, for preparing PRO polypeptides, for generation of antisense and DNA, for preparing PRO polypeptides, for generation of therapeutically useful reagents, as probes and for the generic analysis of individuals with genetic disorders as well as for recombinantly expressing the protein and for chromosome identify contropes, as therapeutic agents, for screening compounds to identify the proteins are useful for tissue typing. PRO antibodies are useful in diagnostic assay of sample fluids. Anti-PRO contibodies are useful in diagnostic assay of sample fluids. Anti-PRO pulyingenes may also be used in gene therapy, particularly for replacing a general part presented is a PCR primer which was used to mplify a pRO polynucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; dabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lery Match 0.4%; Score 14.4; DB 1; Length 18; set Local Similarity 93.8%; Pred. No. 1.7e+03; atches 15; Conservative 0; Mismatches 1: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted/transmembrane protein, #44, PCR primer #2.
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17-SEP-1997;
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9705-0059121P
9705-0059122P
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New isolated nucleic acid molecule for use in molecular biology, as hybridization probe, in chromosome and gene mapping, and in generation of anti-sense ribonucleic acid and deoxyribonucleic acid.
                                                                                                                                                         Botstein D, Desnoyers L, Eaton DL,
Fong S, Gao W, Gerber H, Gerritsen
Grimaldi JC, Gurney AL, Hillan KJ,
an J, Paoni NP, Roy MA, Stewart TA,
                                                                                                                                                                                                                                                                 Example 36; SEQ ID NO 229; 472pp; English.
16-DEC-1999; 99WO-US030095.
20-DEC-1999; 99WO-US0300911.
20-DEC-1999; 99WO-US030991.
11-FEB-2000; 2000WO-US003219.
11-FEB-2000; 2000WO-US003414.
24-FEB-2000; 2000WO-US0034414.
24-FEB-2000; 2000WO-US0050414.
24-FEB-2000; 2000WO-US005041.
20-MAR-2000; 2000WO-US005371.
                                                                                     22-MAY-2000; 2000WO-US014042
02-JUN-2000; 2000WO-US015264
28-JUL-2000; 2000WO-US020710
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18-SEP-2000; 2000US-00665350.
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Godowski PJ, Grim
                                                                                                                                                          Ashkenazi A,
Filvaroff E,
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Ferrara N; NE, Goddard A; Kljavin IJ; Tumas D; The invention discloses isolated PRO secreted/transmembrane polypeptides and the mucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating to the asset one biological activity of a cell. PRO polypeptides are useful for detecting other PRO polypeptides in a sample and for linking a bioactive molecule to a cell expressing a PRO polypeptides are useful for detecting other PRO polypeptides. The PRO polypeptides are useful submardeuticals, diagnostics, biosensors or bioactors. These are useful for stimulating hypertrophy of neonatal proliferation of bioreactors. These are useful for stimulating hypertrophy of neonatal proliferation of cheart, inhibiting vascular endothelial growth factor (VBGF) estimulated proliferation of endothelial growth factor (VBGF) estimulated proliferation of endothelial growth factor (VBGF) estimulated proliferation of chondrocytes, inducing c-fos in endothelial cells, modulating glucose or FRA uptake, inducing proliferation of endothelial cells, inducing proliferation of endothelial growth factor (VBGF) estimulated cumours, retinal disorders or injuries (e.g. loss of sight due to cells, modulating glucose or FRA uptake, inducing proliferation moderolar brone or cartilage disorders (e.g. loss of sight due to cetting or treating cardiac insufficiency disorders (e.g. loss of sight due to cretinists) in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polymucleotides are useful in molecular biology including uses as hybridisation probes or businals or knockout animals which are useful in the development and gene mapping, in the generation of antisenes RNA animals or knockout animals which are useful in the development and gene mapping, in the generation transgenic contening of therapeutically useful reagents, so process as therapeutic aspects and DNA, for preparing PRO polypeptide antibodies are useful in diagnostic assays for PRO e.g. detecting its expression in specific cells, tissues or serum and for affinity purification of PRO from recombinant cell culture or natural sources. The PRO genes may also be used in gene therapy, particularly for replacing a

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tissue typing; immunohistochemical staining; gene therapy; neconatal heart; vascular endothelial growth factor; VEGF; proliferation; endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c-fos; glucose; FFA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinaulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthritis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiarthritic; anorectic.
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defective gene. The sequence presented is a PCR primer which was used amplify a PRO polynucleotide of the invention.
                                                                                     Gaps
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                                                           Query Match
0.4%; Score 14.4; DB 1; Length 18;
Best Local Similarity 93.8%; Pred. No. 1.7e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
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                                    Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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(GETH) GENENTECH INC.

ME, Goddard A; Kljavin IJ; Tumas D; Ferrara N; Botstein D, Desnoyers L, Eaton DL, Fong S, Gao W, Gerber H, Gerritsen Grimaldi JC, Gurney AL, Hillan KJ, an J, Paoni NF, Roy MA, Stewart TA, Gurney AL, F, Roy MA, Wood WI; Ashkenazi A, Botste Filvaroff E, Fong & Godowski PJ, Grima Mather JP, Pan J, Williams PM,

WPI; 2004-096547/10.

Sixty one isolated nucleic acids encoding a PRO polypeptide, e.g. PRO245

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specificially bind to the PRO polypeptide for linking a cariobodies that specificially bind to the PRO polypeptide. The PRO polypeptide of the condulating at least one biological activity of a cell. PRO polypeptides are useful for modulating the biological activity of solypeptides are useful for modulating the biological activity of polypeptides are useful for modulating the biological activity of conducting the biological activity of polypeptides are useful for modulating the biological activity of polypeptides are useful for stimulating hypertrophy of meaners of polymucleotides are useful for stimulating hypertrophy of meaners of polymucleotides are useful for stimulating proliferation of candidating glucose or FRP uptack, inducing proliferation of redinating glucose or FRP uptack, inducing proliferation of chondrocytes. In particular, these are useful for cells, modulating glucose or FRP uptack, inducing proliferation of chondrocytes or inputicis or repeasing cardiac inputies (e.g. loss of sight due to cells, modulating glucose or FRP uptack. Inducing proliferation of chondrocytes or inputicis (e.g. loss of sight due to cells, modulating plucose or FRP uptack. Appring proliferation of chondrocytes or inputicis (e.g. sports injuries or arthritis) in mammals. PRO polypeptides and their portions affect the cerinitis of genes which have a role in cell cells, modulating plucose or cartilage disorders wounds, cancerous tumours, retinal disorders or injudy useful respectives and their portions and provides and problem or cartilage disorders as which have a role in cell for cond library to cisclate the full-length RRO colls, or some proper and publy, for preparing RRO polypeptides, for generating respective provides and provides and publy for preparing RRO polypeptides, for generating cells, inducing seems and publy or crossing the procein and for riseus t .. 0 or PRO1868, useful in chromosome and gene mapping, in generating antisense RNA and DNA, and in treating cancer and Alzheimer's disease. Gaps .. 0 / Match 0.4%; Score 14.4; DB 1; Length 18; Local Similarity 93.8%; Pred. No. 1.7e+03; les 15; Conservative 0; Mismatches 1; Indels Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Example 36; SEQ ID NO 229; 483pp; English. PRO1868, useful in chromosome 1101 GCTGTCCTCAGGGGAG 1116 gererecacaegeae ٣ Query Match Best Loc Matches à 셤

ADM29942 standard; DNA; 18 BP (first entry) 20-MAY-2004 ADM29942; MAKE KE KE KAKA

RESULT 1964

ADM29942

Human secreted/transmembrane protein, #44, PCR primer #2.

Human; PCR; primer; ss; PRO; secreted; transmembrane; therapeutic; tissue typing; immunohistochemical staining; gene therapy; neonatal heart; vascular endothelial growth factor; VEGF; proliferation;

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endothelial cell; stimulated T-lymphocyte; retinal neuron; rod photoreceptor cell; c.fos; glucose; FRA; chondrocyte; cardiac insufficiency disorder; wound; cancer; tumour; retinal disorder; retinitis pigmentosum; obesity; diabete; hyperinsulinaemia; hypoinsulinaemia; bone disorder; cartilage disorder; sport injury; arthitis; cardiant; vulnerary; cytostatic; ophthalmological; osteopathic; antiathritic; anorectic.
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97US-0063564P.
97US-0063435P.
97US-0063704P.
97US-0063732P.
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                                                                         Homo sapiens.
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98WO-US025108.
98US-0113296P.
99US-0143048P.
99US-0145698P.
99WO-US020594.
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                                   99WO-US021090
                                      99WO-US021547
                                         99WO-US023089
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99WO-US028313
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                                                                                    02-MAR-2000;
                                                            16-DEC-1999
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(GETH) GENENTECH INC.

Ferrara N; ME, Goddard A; Kljavin IJ; Desnoyers L, Eaton DL, W, Gerber H, Gerritsen Gurney AL, Hillan KJ, VF, Roy MA, Stewart TA, ldi JC, Gurney AL, Paoni NF, Roy MA, Botstein D, Desr Fong S, Gao W, Grimaldi JC, Gui Mather JP, Pan J, Pac Williams PM, Wood WI; Ashkenazi A, Filvaroff E, Godowski PJ,

WPI; 2004-020978/02.

New PRO nucleic acid, useful for preparing a composition for treating tumor or for tissue typing e.g.,

Example 36; SEQ ID NO 229; 472pp; English

The invention discloses isolated PRO secreted/transmembrane polypeptides and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating a confective molecule to a cell expressing a PRO polypeptides are useful confective molecule to a cell expressing a PRO polypeptides are useful confective molecule to a cell expressing a PRO polypeptide antibodies are useful for modulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptide or collective are useful for stimulating the biological activity of a cell expressing PRO polypeptides. The PRO polypeptides or polymcleotides are useful for stimulating hypertrophy of neonatal collective inhibiting vascular endothelial growth factor (VEGP)-stimulated collection of endothelial cells, modulating the proliferation of collective inhibiting glucose or FPA uptake, inducing coliferation of collection of chondrocytes. In particular, these are useful for cells, modulating proliferation and/or recolls, modulating proliferation and/or recolls, modulating proliferation of chondrocytes. In particular, these are useful for cells, modulating proliferation of chondrocytes. In particular, these are useful for cells, modulating proliferation of chondrocytes. In particular, these are useful for cells, modulating proliferation of chondrocytes. In particular, these are useful for cells, modulating proliferation of chondrocytes. In particular, these are useful for cells, modulating proliferation and/or reconsecting or treating or treating cardiac insufficiency disorders, wounds, cetting the conferons or injuries (e.g. loss of sight due to cetting in mammals. PRO polypeptides and their portions affect the expression of genes which have a role in cell death. The polynuclees or cartilage disorders (e.g. sports injuries or expression of genes which have a role in cell death. The polynuclees

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useful in molecular biology including uses as hybridisation probes
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ADO06264 standard; DNA; 18 RESULT 1965

BP

(first entry) Human PRO PCR primer #102. 01-JUL-2004 AD006264;

Human; PRO; 88; affinity purification; PCR; primer.

Homo sapiens.

US6686451-B1

03-FEB-2004.

10-JUL-2001; 2001US-00902775

24-OCT-1997; 97US-0063128P. 16-SEP-1998; 98WO-US019330. 30-NOV-1999; 99WO-US028113. 22-FEB-2000; 2000WO-US04414. 18-SEP-2000; 2000US-00665350.

(GETH) GENENTECH INC

Mather JP; Godowski PJ, Gurney AL, Goddard A, Wood WI; Desnoyers L, Williams PM,

WPI; 2004-106364/11.

New antibodies binding PRO polypeptides, useful in gene therapy, or in diagnostic assays for the PRO polypeptides, or for the affinity purification of PRO polypeptides from recombinant cell culture or natural

Example 36; SEQ ID NO 229; 445pp; English.

The invention relates to an antibody that binds to a human PRO

The invention describes an isolated polypeptide (I) comprising a cids as given in the specification or a region consisting of 5 or more contiguous amino acids, where the region includes amino acid of 634 of contiguous amino acids, where the region includes amino acid of 634 of thily defined sequence (82) of 2405 base pair as given in the specification, or its complement, a polymucleotide (II) comprising a fully defined sequence (82) of 2405 base pair as given in the specification, or its complement, a polymucleotide that selectively hybridizes to (82) relative to a known polymucleotide, or a region of 15 or more contiguous mucleotides, the region comprising nucleotide 1904 of (82); vector, preferably an expression vector (III) comprising of (11); a chost cell (IV) comprising (II); a leter sample; detecting observe or observer or chatted disorders associated with differential expression of (II); a LIPINN controlled of (IV) or (VI), where the region is 10 or more comprising a detecting a level of expression of (V) or (VI), where the region is 10 or more comprising to (II); an ecombinant cell sample; antibodies that specifically bind to (I); a recombinant cell sample; antibodies that specifically such that the (II) is overexpressed; a composition comprising (I); an array comprising two or more (III); and identifying an alteration in IIPINN gene associated with obesity or an obesity related disorder. (II) ö polypeptide. The invention also relates to human PRO polynucleotides encoding the PRO polypeptides of the invention. The antibody is a monoclonal or humanised antibody, or is an antibody fragment, and is preferably labelled. The anti-PRO antibodies may be used in diagnostic assays for PRO, or for the affinity purification of PRO from recombinant cell culture or natural sources. This sequence represents a PCR primer used in isolation of a human PRO polynucleotide of the invention. Gaps Novel isolated LIPIN3 polypeptide, useful for diagnosing diabetes ö LIPIN3; obesity; obesity-related disorder; differential express polynucleotide polymorphism; adipocyte; human; PCR; primer; ss. 0.4%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 1.7e+03; tive 0; Mismatches 1; Indels Seguence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Human LIPIN3 exon10 PCR primer segid 34. Example 5; SEQ ID NO 33; 58pp; English. 1101 GCTGTCCTCAGGGGAG 1116 ADO17042 standard; DNA; 18 BP 26-JUL-2002; 2002US-00206618. 26-JUL-2002; 2002US-00206618. 3 GCTGTCCACAGGGGAG 18 01-JUL-2004 (first entry) Query Match
Best Local Similarity 93.8
Matches 15; Conservative WPI; 2004-122019/12. (WARD/) WARDEN C H. US2004018497-A1. Homo sapiens. 29-JAN-2004. Warden CH; ADO17042; RESULT 1966 8ઠ 셤

HIV gag homology region and regulatory factor (human RIP protein kinase).

09-MAR-1999 (first entry)

AAV99272;

defibriotide; polyanion salt; HIV; protozoan infection; schistosoma; Schistocerca Leishmania; Trypanasoma; fungus infection; Presuncystis carinii; malaria; viral infection; genetic disease; Duchenne's muscular dystrophy; Down's syndrome; degenerative disease; neoplasia; cancer; skin condition; drug resistance; ss.

Human immunodeficiency virus

Synthetic.

Homo sapiens.

WO9848843-A1

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAQ93237-40 are primers which were used in the amplification of a gene fragment from the acetyl CoA carboxylase (ACCase) gene. The full length ACCase gene may be used to produce a transgenic plant such that it expresses a changed amount of ACCase. The ACCase gene may be used to regulate the production of fat/protein in agricultural
is useful for detecting a polynucleotide polymorphism associated with obesity. (II) is useful for diagnosing obesity an obesity-related disorder which involves detecting (I). In (MZ), the cell is an adipocyte. The test agent is chosen from antibody, protein, nucleic acid, and small organic molecule. This sequence represents a primer used to identify single nucleotide polymorphisms in the human Lipin3 gene that may be associated with obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymerase chain reaction, PCR, primer; amplify, acetyl CoA carboxylase, ACCase, transgenic plant, regulation, fat, protein, agricultural plant,
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Pred. No. 1.8e+03;
1; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 13; 33pp; Japanese.
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Best Local Similarity 83.3%;
Matches 15; Conservative
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AAQ93237/c
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Use of defibrotide nucleic acid components - for treating e.g. infectious diseases, genetic diseases, degenerative diseases, DNA damage, neoplasia and skin disease, particularly HIV infection.

WPI; 1999-034643/03.

Burcoglu A;

(BURC/) BURCOGLU A.

28-APR-1997;

98WO-US008357.

28-APR-1998; 35-NOV-1998

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Contraining a Human immunodeficiency virus (HIV) homology region and a cellular regulatory factor. Defibriotide is a polyanion salt of a decoxythomology region and a cellular regulatory factor. Defibriotide is a polyanion salt of a decoxythomology region and a cellular regulatory factor. Defibriotide is a polyanion salt of a decoxythomology region and a cellular regulatory factor. Defibriotide is a polyanion salt of a decoxythomology of the products can be used for treating diseases such as infection e.g. Schistosoma infection, protozoan infection, schistosoma infection, protozoan infection, schistosoma infection e.g. Schistosoma infection, and fungus infection, pneumocystis carini infection, malaria, plasmodium vivax, gram negative bacterial infection, infection, infection, departitis virus infection, human papilloma virus infection; genetic diseases e.g. buchenne's muscular dystrophy and virus infection; genetic diseases e.g. buchenne's muscular dystrophy and virus infection; genetic diseases e.g. neuropathy, cardiomyopathy, aging, Kearn's Sayre syndrome, retinitis pigmentosa, ataxia, seizures, adiseases, parkinson's disease, neuropathy, cardiomyopathy, aging, Kearn's Sayre syndrome, retinitis pigmentosa, ataxia, seizures, consuritis, and radiation damage; neoplasia, e.g. lympho-proliferative consuritis, and radiation damage; neoplasia, e.g. lympho-proliferative angiomatosis, seborrheic dermatitis, psoriasis, Reiter's syndrome, insect colon cancer; and skin diseases, e.g. molluscum contegiosum, bacillary angiomatosis, seborrheic dermatitis, psoriasis, Reiter's syndrome, insect bite reaction, Staphylococcal folliculitis, Ecosinophilic folliculitis. In addition a drug resistance can be treated via administering the nucleic acid components of defibrotide and the variants in combination with the
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Gaps

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1354 GAGATGATGAAGATGATC 1371

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AAV99272 standard; DNA; 19 BP

RESULT 1968 AAV99272/c ID AAV9927;

(ORCH-) ORCHID BIOSCIENCES INC

99US-0160096P.

15-OCT-1999;

13-OCT-2000; 2000WO-US028436.

WO200129262-A2

26-APR-2001

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CIKI, pcNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                            Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                               New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP specific lower PCR primer SEQ ID 2918.
                                                                                                                     Cyclin E ribozyme binding site #261.
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                              AAA84728 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-412314/35
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Best Local Similarity
Matches 15; Conserv
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                                                                                          04-DEC-2000
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                                                                                                                                                                                                                                             08-JUN-2000
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                                                                                                                                                                                   Mammalia.
                                                            AAA84728,
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Homo sapiens

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performance recovered to the presence for prince of a full control of primers of a full control of primers. The present invention sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the includes kits for determining the presence or absence of a swip using the coligonucleotides of the invention. The PCR primers are used to amplify a coliforning as a ser useful for genotryping a nucleic caid sample by performing a single-nucleotide primer extension reaction. The performing a single-nucleotide primer extension reaction. The cligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotype of an individual or group of sesses by association analysis the genotypic of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. capsembly familial hypercholesterolaemia polycystic kidney disease, dystrophy, familial hypercholesterolaemia, polycystic kidney disease, or steogenesis imperfects and acute intermittent porphyria. Phenotypic traits also include symptoms of or may be genetic such as autoimmune disease of which a component is or may be genetic such as autoimmune inflamment, inflamment, who may never intermitent prompt infertion hyperhorial activities multiple sclerosis, inflamment, and a such as autoimmune inflamment of inflamment in partner and a second second inflamment in the partner and a second second inflamment in the partner of a second inferior and a second second inflamment in the partner of a second 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific
                                                                                                                                                                                                                                                                                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences AAH37205 - AAH40944 represent PCR primers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 3 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            For a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 64; 83pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2736 CCTGAAGATGGGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19 CTTGAAGATGGGAGCC
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                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-290930/30.
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               acid sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                          Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   d hypertrophic or hypertrophic burn
sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human c-fos transcript target sequence/siNA upper strand, SEQ ID NO:40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA interference, short interfering nucleic acid, siNA, short interfering RNA, siRNA, double-stranded RNA, micro-RNA; mirnA, short hairpin RNA, shRNA, expression modulation; gene therapy; day screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
atopic dermatitis, actinic keratosis, squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 14.4; DB 1; Length 19; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     celoid, adhesion and hypertrophic
AAH62099 represent sequences used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                Example 1; Page 240; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2868 TGGTACAACGGAGGCC 2883
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                                                                                                                                             26-OCT-2000; 2000WO-US029500.
                                                                                                                                                                       26-OCT-1999; 99US-0161532P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE65585 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      scarring such as keloid,
scar. AAH57577 to AAH620
                                                                                                                                                                                                                          Robbins JM, Tritz R;
                                                                                                                                                                                                (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                   WPI; 2001-300427/31.
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                                                                                           WO200130362-A2
                                                     Homo sapiens
Synthetic.
                                                                                                                    03-MAY-2001.
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Matches
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human c-fos gene by RNA interference. The siNAs may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense cligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes for the in vitro or in vivo delivery of siNA; conjugates and/or complexes corpuses in calls, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the reatment of a variety of conditions. They may be used for treating central nervous system lesions and injurise (e.g., Alzheimer's disease, Huntington's disease, epilepsy, dementia or myotrophic lateral sclerosis), various cancers; other proliferative and/or allergic diseases; viral infections (including HIV infection);
central nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; epilepsy; dementia; amyotrophic lateral sclenosis; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; viral infection; HIV infection; autoimmune disease; transplant rejection; vasorropic; nocroropic; antiparkinsonian; neuroprotective; cytostatic; antinflammatory; antiallergic; virucide; anti-HIV; immunosupressive; anticonvulsant; nephrotropic; human; c-fos; target sequence; ss.
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targeted double-stranded siNA, which is identical to the c-fos transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid downregulates expression of the c-fos gene useful for treatment and diagnosis of diseases, e.g. cancer and inflammation.
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human c-fos gene by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense algonucleotide and an antisense oilgonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically synthesised, expressed from a coctain decoxyribonucleotide, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits of the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the c-fos gene in cells, tissue explants or organisms of since avariest of conditions. They may be used for treating central nervous system lesions and injuries (e.g., Alzheimer's disease, Extracting central nervous system lesions and injuries (e.g., Alzheimer's disease, Rathinson's disease, Huntington's disease, epilepsy, dementia or amyotrophic lateral sclerosis); various cancers; other proliferative
                                                                                                                                                                                                                                                                       RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mirnA; short hairbin RNA; siRNA; cxpression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; central nervous system disorder; Alzheimer's disease; parkinson; disease; hartington's disease; pliepsy; dementia; amyotrophic lateral sclerosis; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; allergic disease; viral infection; HVI infection; autoimmune disease; transplant resjection; vascircipic; noctropic; antiparkinsonian; neuroprotective; cytostatic; anticonvulsant; nephrotropic; human; c-fos; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New short interfering nucleic acid downregulates expression of the c-fos gene useful for treatment and diagnosis of diseases, e.g. cancer and inflammation.
                                                                                                                                                                                                                                   Human c-fos sinA lower strand, SEQ ID NO:156
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06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409293P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                        BP.
3 CAACCUGCUGAAGGAG 18
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                                                                                                        ADE65701 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beigelman L;
                                                                                                                                                                                              (first entry)
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diseases (e.g., restenosis and polycystic kidney disease); inflammatory and/or allergic diseases; viral infections (including HIV infection); autoimmune diseases; and transplant rejection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene the present sequence represents the lower strand of a human c-fostargeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  short interfering nucleic acid; siNA; dównregulation; inhibition; SCD; stearcyl-CoA desaturase; RNA interference; anorectic; antidiabetic; antiatreiosclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss.
                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                     0.4%; Score 14.4; DB 1; Length 19; 3.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stearoyl-CoA desaturase siNA oligonucleotide SEQ ID NO:262.
                                                                                                                                                                   Sequence 19 BP; 2 A; 6 C; 5 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; SEQ ID NO 262; 139pp; English.
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                                                                                                                                                                                                                                                                                  1887 CAAGCTGCTGAAGGAG 1902
                                                                                                                                                                                                                                                                                                                                                                                                              ADE27318 standard; RNA; 19 BP
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09-SEP-2002; 2002US-0409293P.
20-SEP-2002; 2002US-0412304P.
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29-AUG-2002; 2002US-0406784P.
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                                                                                                                                                                                                                           93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                     17 CAACCTGCTGAAGGAG
                                                                                                                                                                                                                             Local Similarity 93.8 nes 15; Conservative
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                                                                                                                                                                                                            Query Match
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genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target identification and validation; genetic enquineering; pharmacogenomics; studying gene function and spene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    short interfering nucleic acid; siNA; downregulation; inhibition; SCD; stearoyl-COA desaturase; RNA interference; anorectic; antidiabetic; antidiatericsclerotic; cytostatic; virucide; obesity; diabetes; atherosclerosis; cancer; viral infection; drug screening;
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                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.4; DB 1; Length 19;
81.2%; Pred. No. 1.8e+03;
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                                                                                                                                                                                                                                                                                  Sequence 19 BP; 4 A; 5 C; 6 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                     2; Mismatches
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
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15-JAN-2003; 2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                                                                                     13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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diabetes (types I and II); atherosclerosis; cancer and viral infections. They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents an SCD siNA, which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                  1; Indels
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0.4%; Score 14.4; DB 1;
Best Local Similarity 93.8%; Pred, No. 1.8e+03;
Matches 15; Conservative 0; Mismatches 1;
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2002WO-US017674.
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29-JUL-2002; 2002US-0399348P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409293P.
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27-NOV-2002; 2002US-00306747.
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11-MAR-2002;
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endothelial growth factor receptor (VEGFR) gene. Also described: (1) a sink that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of sink; (3) conjugates and/or complexes of sink; (4) vectors that express sink; and (5) single-stranded sink with similar properties. The sinks have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and cativities. The sink are useful for modulating (downregulating) the expression of VEGFR genes. The sink are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The sink may also be useful for diagnosis, drug exceening, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
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Pred. No. 1.8e+03;
0; Mismatches 1; Indels
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29-MAY-2002; 2002WO-US017674.
06-JUN-2002; 2002US-0386782P.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
04-NOV-2002; 2002US-0587949.
27-NOV-2002; 2002US-00306747.
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29-JUL-2002; 2002US-0399348P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2003; 2003WO-US005022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 93.8%;
nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF37445 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1977
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IID ADF3

AC ADF3

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XX

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ADF3

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ADF3

XX

ADF3

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The present invention describes a double-stranded short interfering nucleic acid (sinA) that downregulates expression of the vascular endothelial growth factor receptor (WGGFR) gene. Also described: (1) a caid that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of sinA; (3) conjugates and/or complexes of sinA; (4) vectors that express sinA; and (5) single-stranded sinA with similar properties. The sinAs have antianspicognia, cytostatic, nephrotropic and gynaecological activities. The sinA are useful for modulating commanded sinAs are useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, consovantic kidney disease. The sinA may also be useful for diagnosis, and polycystic kidney disease. The sinA may also be useful for diagnosis, and sories strains, studying gene function, and also for gene mapping (e.g. of single-incleants). The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           short interfering nucleic acid; SiNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; antidiabetic; approximation; and propertions; and gradional and gradional antihities; portable; and degeneration; neovascular glaucoma; polycystic kidney disease; ss.
                                 New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human VEGFR3 short interfering nucleic acid (siNA) SEQ ID NO:1981
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 19;
81.2%; Pred. No. 1.8e+03;
tive 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 6 A; 4 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  double-stranded short interfering nucleic acid;
                                                                                                          Example 3; SEQ ID NO 1734; 207pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                   exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1998 CAAGCAGCTGGTGGAG 2013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002US-0386782P.
2002US-0393796P.
2002US-0399348P.
2002US-0406784P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF37692/c
ID ADF37692 standard; RNA; 19 BP.
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2002WO-US017674.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 CAAGAAGCUGGUGGAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13; Conservative
                                                                          factor receptor gene.
WPI; 2003-679876/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003070910-A2.
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03-JUL-2002;
29-JUL-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAR-2002;
29-MAY-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF37692;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endotherial growth factor receptor (VEGFR) gene. Also described: (1) a sinA that downregulates the VEGF gene. (2) kits for in vitro or in vivo delivery of siNA, (3) conjugates and/or complexes of siNA, (4) vectors that express siNA, and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, angineering, studying gene function, and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of single-nuclectide polymorphisms). The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                         New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 2 A; 7 C; 4 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 1981; 207pp; English.
                                                                                                                                                                                                         Mcswiggen J, Beigelman L, Pavco P;
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
04-NOV-2002; 2002US-00287949.
27-NOV-2002; 2002US-0036747.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                 factor receptor gene.
                                                                                                                                                                                                                                                            WPI; 2003-679876/64.
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; 0 0.4%; Score 14.4; DB 1; Length 19; 13.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels 93.8%; Query Match
Best Local Similarity 93.8%

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Gaps

1998 CAAGCAGCTGGTGGAG 2013 17 CAAGAAGCTGGTGGAG 2 ઠ 셤

ADF36753 standard; RNA; 19 BP. ADF36753; RESULT 1979 ADF36753
IID ADF3
XX AC ADF2
XXX
XX I2-E
XXX
XX GOUL
XXX
XX GOUL
XXX
XX GOUL
X

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12-FEB-2004 (first entry)

Human VEGFR2 short interfering nucleic acid (siNA) SEQ ID NO:1042.

double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopsthy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; ss.

Homo sapiens. Synthetic.

WO2003070910-A2

28-AUG-2003

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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene, (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antianglogenic, cytostatic, antidabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of engine-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                       New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; Myc; Myb; cancer; proliferative disease; restenosis; polycystic kidney disease; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA, adouble-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis;
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31.2%; Pred. No. 1.8e+03;
ve 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 6 A; 4 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; SEQ ID NO 1042; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human Myb siNA lower strand, SEQ ID 572.
                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J, Beigelman L, Pavco P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1609 AAGTGCATCCACAGGG 1624
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                                                                                                                                                                                              05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
04-NOV-2002; 2002US-00287949.
                                                                                                                 2002US-0386782P
2002US-0393796P
                                                                                                                                                        2002US-0399348P
                                                                                                                                                                                                                                                        27-NOV-2002; 2002US-00306747
15-JAN-2003; 2003US-0440129P
                  20-FEB-2003; 2003WO-US005022
                                                                                                 2002WO-US017674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 AAGUGUAUCCACAGGG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 81.2%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-679876/64.
                                                                                                                 06-JUN-2002;
03-JUL-2002;
                                                                                                                                                           29-JUL-2002;
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                                                                                             29-MAY-2002;
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MANA MAKA BAKA KA KA D
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Human Myb transcript target sequence/siNA upper strand, SEQ ID 393

Human; Myc; Myb; cancer; proliferative disease; restenosis; polycystic kidney disease; RNA interference; short interfering RNA; siRN double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cytostatic; vasotropic;

nephrotropic; ss.

WO2003070917-A2.

28-AUG-2003.

Homo sapiens.

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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of Myc or Myb genes
therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cytostatic; vasotropic; nephrotropic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 11 A; 0 C; 0 G; 0 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                  Example 7; Page 135; 161pp; English
                                                                                                                        11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
                                                                                                                                                   05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-OCT-2002; 2002US-041655EP.
15-JAN-2003; 2003US-0440129P.
                                                                                               20-FEB-2003; 2003WO-US005326
                                                                                                                2002US-0358580P
                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                   Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                   WPI; 2003-689784/65.
                                                          WO2003070917-A2.
                                         Homo sapiens.
                                                                                                               20-FEB-2002;
                                                                            8-AUG-2003
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human Myc or Myb genes by RNA interference. The siNAs may or may not comprise ribonuclecides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclecide and antisense oligonuclecide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the Myc or Myb genes in cells, tissue caplants for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the Myc or Myb genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancers and other proliferative diseases, such as restenosis and polycystic kidney disease. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human Myb-targeted double-stranded siNA.

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Gaps
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0.4%; Score 14.4; DB 1; Length 19; 50.0%; Pred. No. 1.8e+03; ve 7; Mismatches 1; Indels
               50.08;
               Local Similarity 50.0
hes 8; Conservative
   Query Match
                                 Matches
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(first entry) 12-FEB-2004 RESULT 1981
ADF48256/c
ID ADF48256
XX
AC ADF4825
XX
DT 12-FEB-3

ADF48256;

ADF48256 standard; RNA; 19 BP

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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of Myc or Myb genes

11-MAR-2002; 2002US-0363124P. 06-UTV-2002; 2002US-036782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 15-CT-2002; 2002US-040923P. 15-JAN-2003; 2003US-0418655P.

(RIBO-) RIBOZYME PHARM INC

Beigelman L;

Mcswiggen J,

WPI; 2003-689784/65.

20-FEB-2003; 2003WO-US005326

Example 7; Page 135; 161pp; English.

seguence ö

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human Myc or Myb genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense of cregions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA, and vectors that express siNA. The siNAs are and/or complexes of siNA, and vectors that express siNA. The siNAs are captaints or organisms (e.g., by ex vivo gene therapy), or in grafts and captaints for the treatment of a variety of conditions. They may be consed for modulates expression of the Myc or Myb genes in cells, tissue captaints for the treatment of a variety of conditions. They may be consed for treating cancers and other proliferative diseases, such as createness and polycystic kidney disease. The siNAs are also useful for drug screening, denetic engineering, pharmacogenomics, studying gene containts sequence represents the upper strand of a human Myb-targeted containers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 8 A; 0 C; 0 G; 0 T; 11 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2833 TATATATATACAT 2848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19 rararararaaar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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ss; siNA; human; BCL2; short interfering nucleic acid; RNA interference; cytostatic; immunosuppressive; virucide; anti-HIV; cancer; autoimmune disease; viral infection; HIV.

18-FEB-2003; 2003WO-US004908

WO2003070969-A2 Homo sapiens.

28-AUG-2003.

Human BCL2 siNA upper sequence SEQ ID NO:543

(first entry)

12-FEB-2004

ADF49815;

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sinA of the invention has cytostatic, immunosuppressive, virucide, and anti-HIV activity. The sinA are useful for modulation (inhibition) of expression or activity of BCL2 by RNA interference. sinA are used to modulate expression of BCL2 genes, in cells, tissue explants or organisms, e.g. for treating cancer, autoimmune diseases and viral infections (including by HIV) but also for drug screening, diagnosis, pharmacogenomics, genetic engineering, pharmacogenomics, studying gene function and gene mapping e.g. of single-nucleotide polymorphisms). The sequences shown in ADF49273-ADF50143 represent sinA of the invention.
                                                                                                                                                                                                                                                                                                                                                                            ss; siNA; human; BCL2; short interfering nucleic acid; RNA interference; cytostatic; immunosuppressive; virucide; anti-HIV; cancer; autoimmune disease; viral infection; HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer or autoimmune disease, downregulates expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel short interfering nucleic acid (siNA) that downregulates expression of the BCL2 gene by RNA interference. A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.4%; Score 14.4; DB 1; Length 19; Best Local Similarity 93.8%; Pred. No. 1.8e+03; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 9 A; 7 C; 2 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                             Human BCL2 siNA upper sequence SEQ ID NO:129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; SEQ ID NO 129; 148pp; English.
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                                                                                  ADF49401 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409233P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
18-JUL-2002; 2002US-0396905P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-712622/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003070969-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                               12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-AUG-2003
                                                                                                                                                        ADF49401;
RESULT 1982
ADF49401/C
XX
AC ADF49401/C
DT 12-FEB-:
XX
BS; SINI
KW CYCOSTS
KW CYCOST
KW CYCOSTS
KW CYCOST
KW CYCOST
KW CY
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The invention relates to a novel short interfering nucleic acid (siNA) that downregulates expression of the BCL2 gene by RNA interference. A siNA of the invention has cytostatic, immunosuppressive, virucide, and anti-HIV activity. The siNA are useful for modulation (inhibition) of expression or activity of BCL2 by RNA interference. SiNA are used to modulate expression of BCL2 genes, in cells, tissue explants or organisms, e.g. for treating cancer, autoimmune diseases and viral infections (including by HIV) but also for drug screening, diagnosis, traget identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The sequences shown in ADF49273-ADF50143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer or autoimmune disease, downregulates expression of the BCL2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein tyrosine phosphatase type IV (PRL3) gene siNA, SEQ ID No 138.
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13.8%; Pred. No. 1.8e+03;
ive 8; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 1 A; 2 C; 7 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 543; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -nucleotide polymorphisms). The represent siNA of the invention.
                                                                                                                                                                                                                                                                           20-PEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
19-UUL-2002; 2002US-0396905P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2316 TCTGTGTGTGTGTG 2331
                                                                                                                                                                                                                                                                                                                                                                                         15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.4*;
Best Local Similarity 43.8%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF71353 standard; RNA; 19
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AC ADF7
XX
DT 12-F
XX
DE Prot
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Gaps

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ADF49815 standard; RNA; 19 BP.

RESULT 1983 ADF49815 ID ADF4

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Homo sapiens.

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short interfering nucleic acid; siNA;
protein tyrosine phosphatase type IV; PRL3; RNA interference; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel short interfering nucleic acid (siNA) that downregulates expression of a protein tyrosine phosphatase type IV (PRL3) gene by RNA interference. The invention further relates to modulating the expression of PRL3 genes in cells, tissue explants or organisms by the introduction of an siNA, kits for in vitro or in vivo delivery of an siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The novel siNA's of the invention have cytostatic activity. siNA's are used to modulate expression of PRL3 genes, in cells, tissue explants or organisms, e.g. for treating cancer but also for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). This polymucleotide sequence represents a short interfering nucleic acid for downregulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    short interfering nucleic acid; siNA; prL3; RNA interference; cytostatic; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression of a protein tyrosine phosphatase type IV (PRL3) gene of the
                                                                                                                                                                                                                                                                                                                                                                                                                                             New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer , downregulates expression of a protein tyrosine phosphatase type IVa gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 138; 131pp; English
                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen J, Beigelman L, Usman N;
                                                                                                                                                                                                    20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                  WO2003070886-A2
                                                                    Homo sapiens
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                                                                                                                                    28-AUG-2003
                                 cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention
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The invention relates to a novel short interfering nucleic acid (siNA)
that downregulates expression of a protein tyrosine phosphatase type IV
(PRL3) gene by RNA interference. The invention further relates to
modulating the expression of PRL3 genes in cells, tissue explants or
organisms by the introduction of an siNA; kits for in vitro or in vivo
delivery of an siNA; conjuages and/or complexes of siNA; and vectors
that express siNA. The novel siNA; of the invention have cytostatic
activity. siNA's are used to modulate expression of PRL3 genes, in cells,
tissue explants or organisms, e.g. for treating cancer but also for drug
screening; diagnosis; target identification and validation; genetic
engineering; pharmacogenomics; studying gene function and gene mapping
(e.g. of single-nucleotide polymorphisms). This polymucleotide sequence
represents a short interfering nucleic acid for downregulating the
expression of a protein tyrosine phosphatase type IV (PRL3) gene of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human GAB2 short interfering nucleic acid upper sequence SEQ ID NO:292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer , downregulates expression of a protein tyrosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 64; 131pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Usman N;
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                                                                                                                                                                                                      11-FEB-2003; 2003WO-US004347.
                                                                                                                                                                                                                                                                                                                   2002US-0363124P.
                                                                                                                                                                                                                                                                                                                                                2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                    29-AUG-2002; 2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen J, Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphatase type IVa gene.
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les 14; Conservative
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                                                                    WO2003070886-A2
                                                                                                                                                                                                                                                                                                                                                06-JUN-2002;
                                                                                                                                                                                                                                                                           20-FEB-2002;
11-MAR-2002;
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Human GAB2 short interfering nucleic acid lower sequence SEQ ID NO:628

(first entry)

12-FEB-2004

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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the GRB2-associated
                                                                                                                                                                                                                Example 3; SEQ ID NO 292; 140pp; English.
                                                                                                                                                     Mcswiggen J, Beigelman L, Usman N;
                                                                                11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409293P.
                                                           18-FEB-2003; 2003WO-US004909
                                                                                                                       L5-JAN-2003; 2003US-0440129P
                                                                                                                                      (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                           diagnosis of cancer, binding protein gene.
                                                                                                                                                                   WPI; 2003-697611/66.
                             WO2003070903-A2.
               Homo sapiens.
                                                                          20-FEB-2002;
11-MAR-2002;
                                            28-AUG-2003
       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                     sequence.
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The present invention relates to short interfering nucleic acids (siNA)
which downregulate expression of the human GRB2-associated binding
comprise in (GAB2) gene by RNA interference. The siNAs may or may not
comprise ribonucleotides and may be double or single stranded. They
further comprise sense and antisense regions, or alternatively are
ssembled from a sense oligonucleotide and an antisense oligonucleotide.
Specifically, the siNAs include short interfering RNA (siRNA), double-
stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs
can be unmodified or chemically modified, can contain
deoxyribonucleotides, and can be chemically synthesised from a
vector or enzymatically synthesised. The invention also relates to kits
correct or enzymatically synthesised. The invention also relates to kits
for the in vitro or in vivo delivery of siNA; conjugates and/or complexes
of siNA; and vectors that express siNA. The siNAs are used to modulate
expression of the GAB2 gene in cells, tissue explants or organisms (e.g.,
by ex vivo gene therapy), or in grafts and transplants for the treatment
of a variety of conditions. They may be used for treating cancer,
inflammation and allergies. The siNAs are also useful for drug screening,
diagnosis, therapeutic target identification and validation, genetic
engineering, pharmacogenomics studying gene function, and gene mapping
(e.g., of single nucleotide polymorphisms). The human GAB2 tergeted
controm or hromosome 11, more specifically to region 11q13.4 The human
GAB2 sinNa have cytostatic, antinflammatory and antiallergic activities.
The present sequence represents the upper strand of a human GAB2-targeted
double-stranded sinN, which is identical to the GAB2 transcript target
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0.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indels
                                                                            1992 CACCTTCAAGCAGCTG 2007
                                                                                                                19 CCCCTTCAAGCAGCTG
                   Local Similarity 93.8 es 15, Conservative
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The present invention relates to short interfering nucleic acids (siNA)
which downregulate expression of the human GRB2-associated binding
comprise infoRAE) gene by RNA interference. The siNAE may or may not
comprise ribonuclectides and may be double or single stranded. They
further comprise sense and antisense regions, or alternatively are
comprise sense and antisense regions, or alternatively are
further comprise sense and antisense regions, or alternatively are
comprise sense and antisense regions, or alternatively are
stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA), double-
stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAE
compositionally synthesised, can contain
deoxyribonuclectides, and can be chemically synthesised, expressed from a
vector or enzymatically synthesised. The invention also relates to kits
corporate in vitro or in vivo delivery of siNA; conjugates and/or complexes
of siNA; and vectors that express siNA. The siNAS are used to modulate
expression of the GAB2 gene in cells, tissue explants for the treatment
complexed to modulate to a since the complexes of a variety of conditions. They way be used for treating cancer,
inflammation and allergies. The siNAS are also useful for drug screening,
diagnosis, therapeutic target identification and validation, genetic
complexed on chromosome 11, more specifically to region 11q13.4. The human
complexed on chromosome 11, more specifically to region 11q13.4. The human
complexed on thromosome 11, more specifically to region 11q13.4. The human
complexed on thromosome 11, more specifically to region 11q13.4. The human
complexed to the context of the conte
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ADF54555 standard; RNA; 19 BP

Matches

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ADF54555; RESULT 1987
ADP54555
ID ADP54555
XX
AC ADP54555

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; human; gRB2-associated binding protein; GAB2; cancer; inflammation; allergy; chromosome 11; cytostatic; antinflammatory; antiallergic; ss. New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the GRB2-associated binding protein gene. 0.4%; Score 14.4; DB 1; Length 19; 75.0%; Pred. No. 1.8e+03; Live 3; Mismatches 1; Indels Example 3; SEQ ID NO 628; 140pp; English. Mcswiggen J, Beigelman L, Usman N; 1992 CACCTICAAGCAGCIG 2007 2002US-0386782P. 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. 18-FEB-2003; 2003WO-US004909 2002US-0363124P (RIBO-) RIBOZYME PHARM INC Query Match
Best Local Similarity 75.0%
Local 12; Conservative WPI; 2003-697611/66. WO2003070903-A2 Homo sapiens. 29-AUG-2002; 20-FEB-2002; 06-JUN-2002; 11-MAR-2002; 28-AUG-2003 Synthetic ઠે

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates concluded expression of the telomerase genes in calls, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases.
                                                                                                                                                                                                                                                                          Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression medulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                           Human TERT transcript target sequence/siNA upper strand, SEQ ID 57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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11-MAZ-2002; 2002US-0363124P.
06-UIN-2002; 2002US-0386782P.
17-UIL-2002; 2002US-0396600P.
29-AUC-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0406784P.
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                                                                                                                      ADF93340 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                               ADF93340;
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e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence

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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; carression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                      Gaps
represents the upper strand of a human TERT-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                    Score 14.4; DB 1; Length 19;
Pred. No. 1.8e+03;
1; Mismatches 1; Indels
                                                       Sequence 19 BP; 1 A; 3 C; 13 G; 0 T; 2 U; 0 Other;
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06-UTW-2002; 2002US-0386782P.

17-UTL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0408378P.

05-SEP-2002; 2002US-040923P.

15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                            ADF93594 standard; RNA; 19 BP
                                                                                                0.4%;
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                                                                                                                                      14; Conservative
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                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-2004
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                                                                                                                                                                                                                                                                        RESULT 1989
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                                                                                                                                      Matches
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                                                                                                                          protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nuclectide polymorphisms). The present sequence represents the lower strand of a human TERT-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Selecting genetic markers as targets for nucleic acid sequence amplification, useful for improving genetic testing, e.g. fetal sex determination, comprises selecting each of the genetic markers according to a heterozygosity index.
used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 2 A; 13 C; 3 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 14.4; DB 1;
93.8%; Pred. No. 1.8e+03;
tive 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCCGGGCGTGGGGG 2935
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACA88903 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15; Conservative
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The invention describes a method of selecting genetic markers as targets for nucleic acid sequence amplification comprising selecting each of the genetic markers according to a heterozygosity index of 0.5 or greater. Selecting and amplification of genetic markers are useful as targets for nucleic acid sequence amplification, for genetic testing or facilitating multiplex PCR amplification from limiting amounts of target nucleic acid. The methods are also useful for improving genetic diagnostic and screening methods, such as prenatal diagnostic testing, foetal sex determination or genetic identification, e.g. DNA profiling or DNA fingerprinting. The nucleic acid sequence amplification is also useful in

Claim 36; Page 39; 64pp; English.

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The present invention relates to a method of determining a genotype associated with increased or decreased resistance to familial bipolar affective disorder. The method comprises determining the genotype with at least one marker of at least one chromosomal region linked to a locus associated with resistance to bipolar affective disorder, where the chromosomal regions are included of and localised between D45402 and D45404, or D118394 and D11829. The invention also disoloses a kit for determining a genotype associated with increased or decreased resistance to familial bipolar affective disorder, where the ckt comprises markers for two or more of the chromosomal regions cited. The method and kit are useful for determining a genotype associated with increased or decreased resistance to familial bipolar affective disorder in a family affected by bipolar affective disorder contribution of these chromosomal regions to bipolar affective disorder in an affective family member, and for assessing an increased or decreased risk of developing bipolar illness for a tested individual from the research invantion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, genotype determination, familial bipolar affective disorder, chromosomal region linked; locus associated with resistance; D4S402; D4S424; D4S431; D4S404; D11S394; D1S29; chromosome marker; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human familial bipolar affective disorder chromsome marker primer #230.
                                                                                                                                                          Gaps
forensic analysis of degraded, old, ancient and difficult samples are difficult to amplify and identify. This sequence represents a primer used in the selection and amplification of genetic markers
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                                                                                                                0.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indels
                                                                              Sequence 19 BP; 2 A; 2 C; 8 G; 7 T; 0 U; 0 Other;
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97US-0062924P.
98US-00175158.
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(EGEL/) EGELAND J A.
(PAUL/) PAUL S M.
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                                                                                                                                      Local Similarity
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20-OCT-1997;
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ACA58282/
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modulate the expression of a HBV target gene in a cell, tissue or
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24-OCT-2000;
08-JUN-2001;
08-JUN-2001;
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26-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                          ADL99963;
                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                   Virucide, Hepatotropic, Gene therapy, ss, short interfering nucleic acid, siNA, hepatitis B virus; HBV; RNA interference.
                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for characterizing gene function.
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                               Length 19;
                                                     1; Indels
           Sequence 19 BP; 2 A; 2 C; 8 G; 7 T; 0 U; 0 Other;
                               Score 14.4; DB 1;
Pred. No. 1.8e+03;
                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beigelman L;
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                                                                          2101 GACACCCCCAGCTCCA 2116
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08-NOV-1999; 99US-00436430.
20-AMR-2000; 2000US-00531025.
09-AUG-2000; 2000US-00636385.
24-OCT-2000; 2000US-00696347.
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08-JUN-2001; 2001US-02968776P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-03370559P.
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11-MAR-2002; 2002US-0363124P.
26-MAR-2002; 2002WO-US009187.
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29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
                                0.4%;
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                                                                                                                                                  ADL99953 standard; RNA; 19
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                                                                                              19 GACATCCCCAGCTCCA
                                                     15; Conservative
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                                          Local Similarity
                                                                                                                                                                                                                                                                    Hepatitis B virus.
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                                                                                                                                                                       ADL99953;
                                 Query Match
                                                                                                                               RESULT 1992
                                                     Matches
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monutage the expression of a new carget gate in a cell, tisher or monitare the expression of a new corganism; and (v) a method for synthesising a sink molecule. The sink molecule is adapted for use to treat HBV infection, and comprises a sense and an antisense region, where the antisense region comprises sequence encoding HBV and the sense region complementary to the antisense region. The sink molecule is assembled from 2 nucleic acid fragments, where one fragment comprises the antisense region of the sink molecule, where sense region and the antisense region comprise separate oligonucleotides, and are covalently connected via a linker molecule. The linker molecule is a polynucleotide linker or a non-nucleotide linker. The sense region comprises a 3'-terminal overhang and the antisense region comprises a 3'-terminal overhang are region such the sense region comprises a 3'-terminal overhang is complementary to RNA encoding HBV. The sink is useful for treating human hepatitis B virus infections, and for characterising pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function of any also be used in clinical, industrial, environmental, agricultural and or the function analysis. The sink molecules and or research settings. The present sequence represents 1 of 1504 HBV
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99US-00436430.
2000US-00531025.
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2000US-00696347.
2001US-00877478.
2001US-0296876P.
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2001US-0337055P.
2002US-0358580P.
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2002WO-US009187.
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les 10; Conservative
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The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA conterference or that inhibite HBV replication. Also disclosed are the following: (i) a method of modulating the expression of a HBV gene in a tissue explant; (ii) a method of generating a library of siNA constructs complexity; (iii) a cell containing one or more siNA molecules; (iv)a kit containing a siNA molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or conjanism, and (v) a method for synthesising a siNA molecule. The siNA molecule is adapted for use to treat HBV infection, and comprises a sense conjementary to a method for synthesising a siNA molecule. The siNA complementary to an RNA sequence encoding HBV and the sense region comprises sequence complementary to the antisense region comprises the molecule is assembled from 2 nucleic acid fragment comprises the complementary to the antisense region and the second fragment comprises the antisense region of the siNA molecule. The linker molecule is a polynucleotide intisense region of the siNA molecule. The linker molecule is a polynucleotide contisense region of the siNA molecule. The linker molecule is a polynucleotide contisense region of the siNA molecule. The linker molecule is a polynucleotide contisense region of the siNA molecule. The linker molecule is a polynucleotide contisense region of the siNA is useful for treating benefity to determine the function of uncharacterised genes in a pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function continue and vernal surface in gene function analysis. The siNA molecules in an action may also be used in clinical, industrial, environmental, agricultural condition of the invantion environmental of the invantion environment
                                                                                                                                                                                                                          New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for
                                                                                                                Morrissey D, Mcswiggen JA, Beigelman L;
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                                                                                                                                                                                                                                                                                                                  characterizing gene function.
MORRISSEY D.
MCSWIGGEN J A.
BEIGELMAN L.
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  (MORR/)
                                                          (BEIG/)
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ö 0.4%; Score 14.4; DB 1; Length 19; 22.5%; Pred. No. 1.8e+03; ve 5; Mismatches 1; Indels Sequence 19 BP; 5 A; 6 C; 3 G; 0 T; 5 U; 0 Other; 62.5%; Query Match
Best Local Similarity 62.54
These 10; Conservative

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Gaps

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ADM00610 standard; RNA; 19 BP. ADM00610; RESULT 1994

20-MAY-2004 (first entry)

Hepatitis B virus short interfering nucleic acid (siNA) #1026.

Virucide; Hepatotropic; Gene therapy; ss; short interfering nucleic acid; siNA; hepatitis B virus; HBV; RNA interference.

Hepatitis B virus.

US2003206887-A1

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2002WO-US009187.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                             2000US-00531025.
2000US-00636385.
2000US-00696347.
                                                   2001US-0335059P.
2001US-0337055P.
                                                            2002US-0358580P.
2002US-0363124P.
                     94US-00193627.
99US-00436430.
                                          2001US-00877478,
                                                2001US-0296876P.
        2002US-00244647
                                                                                      39-SEP-2002; 2002US-0409293P
        16-SEP-2002;
                                           08-JUN-2001;
                                               08-JUN-2001;
24-OCT-2001;
                                                            20-FEB-2002;
                                                        05-DEC-2001;
                                                                         06-JUN-2002;
                                   09-AUG-2000;
06-NOV-2003
RX
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Morrissey D, Mcswiggen JA, Beigelman L;

(MORR/) MORRISSEY D. (MCSW/) MCSWIGGEN J A. BEIGELMAN L.

BEIG/)

WPI; 2003-901032/82.

New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for characterizing gene function.

Claim 11; Page 46; 72pp; English

The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA interference or that inhibits HBV replication. Also disclosed are the following: (1) a method of modulating the expression of a HBV gene in a tissue explant; (ii)) a method of generating a library of siNA constructs complexity; (iii) a cell containing one or more siNA molecules; (iv)a kit containing a siNA molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or complementary to an RNA sequence encoding HBV and the sense region of an an antisense region, where the antisense region comprises a sense and an antisense region, where the antisense region. The siNA conscrule is assembled from 2 nucleic acid fragments, where one fragment comprises the sense region and the second fragment engines region of the siNA molecule. Where sense region and the comprises the comprises the single region comprise separate oligonucleotides, and are covalently comprises region comprise separate oligonucleotides, and are covalently conformant and the sense region and the conformation and The antisense region 3'-terminal overhang is complementary to RNA encoding HBV. The siNA is useful for treating human hepatitis B virus infections, and for characterising pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function of uncharacterised genes in gene function analysis. The siNA molecules may also be used in clinical, industrial, environmental, agricultural and/or research settings. The present sequence represents 1 of 1504 HBV siNA molecules of the invention.

Sequence 19 BP; 5 A; 3 C; 6 G; 0 T; 5 U; 0 Other;

Gaps ; 0 0.4%; Score 14.4; DB 1; Length 19; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels 93.88; Best Local Similarity 93.8 Matches 15; Conservative Query Match

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The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA interference or that inhibits HBV replication. Also disclosed are the following: (i) a method of modulating the expression of a HBV gene in a tissue explant; (ii) a method of generating a library of siNa constructs having predetermined complexity; (iii) a cell containing one or more siNa molecules; (iv) a kit containing a siNa molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or corganism; and (v) a method for synthesising a siNa molecule. The siNa molecule is adapted for use to treat HBV infection, and comprises a sense and an antisense region, where the antisense region comprises sequence complementary to an RNA sequence encoding HBV and the sense region comprises sequence complementary to the antisense region. The siNA molecule is assembled from 2 nucleic acid fragments, where one fragment
                                                                                                                                                                                                                                                       Virucide, Hepatotropic, Gene therapy, ss, short interfering nucleic acid, siNA, hepatitis B virus, HBV, RNA interference.
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                                                                                                                                                                                                                        Hepatitis B virus short interfering nucleic acid (siNA) #1057.
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07-FEB-1994; 94US-00193627.
20-MAR-2000; 2000US-00531025.
09-AUG-2000; 2000US-00531035.
24-OCT-2000; 2000US-00696347.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0296876P.
25-DEC-2001; 2001US-035059P.
20-DEC-2001; 2001US-035059P.
11-MAR-2002; 2002US-0358560P.
2776 TTCCGGAAACTAGTGT 2791
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                                                                                                                    ADM00641 standard; RNA; 19
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(MCSW/) MCSWIGGEN J A.
(BEIG/) BEIGELMAN L.
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35-SEP-2002;
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comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, where sense region and the cantisense region comprises separate oligonuclocides, and are covalently connected via a linker molecule. The linker molecule is a polynuclectide linker or a non-nucleotide linker. The sense region comprises a 3'-terminal overhang and the antisense region comprises a 3'-terminal overhang or antisense region comprises a 3'-terminal overhang or antisense region of pour 2 nucleotides. The antisense region 3'-terminal overhangs each comprise about 2 nucleotides. The antisense region 3'-terminal overhangs is complementary to RNA encoding HBV. The siNA is useful for treating human hepatitis B virus infections, and for characterising pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function of uncharacterised genes in gene function analysis. The siNA molecules and/or research settings. The present sequence represents 1 of 1504 HBV is INA molecules of the invention.
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20-MAR-2000; 2000US-00531025.
09-MAC-2000; 2000US-0053885.
24-0CT-2000; 2000US-0069347.
08-JUN-2001; 2001US-00877478.
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11-MAR-2002; 2002US-0363124P.
26-MAR-2002; 2002WO-US009187.
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2002US-0406784P
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MCSW/) MCSWIGGEN J A.
(BEIG/) BEIGELMAN L.
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                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis B virus.
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ADL99998
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New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for
                                                                                                                                                                                           Claim 11; Page 46; 72pp; English
                                                                                                                           characterizing gene function.
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The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA interference or that inhibits HBV replication. Also disclosed are the following: (1) a method of modulating the expression of a HBV gene in a tissue explant; (ii)) a method of generating a library of siNA constructs having predetermined complexity; (iii) a cell containing one or more siNA molecules; (iv)a kit containing a siNA molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or organism; and (v) a method for synthesising a siNA molecule. The siNA molecule is adapted for use to treat HBV infection, and comprises sequence complementary to an RNA sequence encoding HBV and the sense region comprises sequence complementary to the antisense region. The siNA molecule is assembled from 2 nucleic acid fragments, where one fragment comprises the sense region and the second fragment comprises the antisense region of the siNA molecule, where sense region and the contribers and an alinker molecule. The linker molecule is a polynucleotide connected via a linker molecule. The linker molecule is a polynucleotide linker or a non-nucleotide linker. The sense region comprises a 3'-terminal overhang and the antisense region or the pNA for comprise and the antisense region comprises a 3'-terminal overhang and the antisense region comprises by the sum antisense region and the antisense region comprises by the pNA antisense region and the antisense region comprises a 3'-terminal overhang and the antisense region comprises a 3'-terminal overhang and the antisense region comprises to bNA antisense region and the antisense region comprises to bNA antisense region and the antisense region comprises a 3'-terminal overhang and the antisense region comprises to bNA antisense region a The antisense region 3.-terminal overhang is complementary to RNA encoding HBV. The siNA is useful for treating human hepatitis B virus infections, and for characterising pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function of uncharacterised genes in gene function analysis. The siNA molecules may also be used in clinical, industrial, environmental, agricultural and/or research settings. The present sequence represents 1 of 1504 HBV siNA molecules of the invention.

Sequence 19 BP; 4 A; 5 C; 4 G; 0 T; 6 U; 0 Other;

ö Gaps ö 0.4%; Score 14.4; DB 1; Length 19; 62.5%; Pred. No. 1.8e+03; tive 5; Mismatches 1; Indels 2776 TTCCGGAAACTAGTGT 2791 Best Local Similarity 62.5 Matches 10; Conservative

UCCGGAAACUACUGU 17

Hepatitis B virus short interfering nucleic acid (siNA) #1061.

Virucide; Hepatotropic; Gene therapy; ss; short interfering nucleic acid; siNA; hepatitis B virus; HBV; RNA interference.

US2003206887-A1.

06-NOV-2003.

16-SEP-2002; 2002US-00244647

92US-00882712 94US-00193627. 14-MAY-1992; 07-FEB-1994; 08-NOV-1999;

08-JUN-2001; 2001US-00877478. 08-JUN-2001; 2001US-0296876P. 24-GCT-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P. 10-PRE-2002; 2002US-0358580P. 11-PAR-2002; 2002US-035324P. 26-WAR-2002; 2002WG-03609187. 06-JUN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P.

MORRISSEY D. MCSWIGGEN J A. BEIGELMAN L. (BEIG/) (MORR/)

Morrissey D, Mcswiggen JA, Beigelman L;

WPI; 2003-901032/82.

New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for characterizing gene function.

Claim 11; Page 46; 72pp; English.

The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA conterference or that inhibits HBV replication. Also disclosed are the following: (i) a method of modulating the expression of a HBV gene in a certification of a HBV gene in a cell containing on or more siNA molecules; (iv) a kit containing a siNA molecule which can be used to modulate the expression of a HBV target gene in a cell, tissue or conjection; and an antisense region of a HBV target gene in a cell, tissue or conjective; (iv) a kit containing a siNA molecule which can be used to conjective; (iv) a kit containing a siNA molecule which can be used to conjective; (iv) a kit containing a siNA molecule which can be used to comprise sequence complementary to the antisense region comprises a sense comprises region, where the antisense region comprises sequence complementary to the antisense region comprises the sense region and the sense region and the comprises the sense region and the sense region of the siNA molecule, where sense region and the connected via a linker molecule. The linker molecule is a polynucleotide inthem the sense region comprises a 3'-terminal overhang and the antisense region comprises a 3'-terminal coverhang and the antisense region comprises a conjection, and for characterising pathways of gene function, e.g. to inflections, and for characterising pathways of gene function, e.g. to inflictions, and for characterising pathways of gene function, e.g. to inflict activity of target genes in a pathway to determine the function of undial activity of target genes in a pathway to determine the function and/or research settings. The sink molecule in the man the present sequence represents 1 of 1504 HBV with a sund or research settings. sinA molecules of the invention.

Sequence 19 BP; 6 A; 4 C; 5 G; 0 T; 4 U; 0 Other;

Gaps ö 0.4%; Score 14.4; DB 1; Length 19; 3.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels 93.8%; Query Match 0.49 Best Local Similarity 93.89 Matches 15; Conservative

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<u>ج</u>ڙ. 셤 RESULT 1998

ADL99994

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(first entry)

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New short interfering nucleic acid molecules which down-regulates expression of a hepatitis B virus (HBV) or which inhibits HBV replication, useful for treating human HBV infections or for
                                                  Hepatitis B virus short interfering nucleic acid (siNA) #411.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 11; Page 46; 72pp; English
                                                                                                                                                                       14-MAY-1992; 92US-00882712.
07-FEB-1994; 94US-00193627.
20-MAR-2000; 2000US-00531025.
09-AUG-2000; 2000US-00531025.
09-AUG-2000; 2000US-00636385.
24-OCT-2000; 200US-00696347.
08-JUN-2001; 200US-00596747.
24-OCT-2001; 200US-0296876P.
24-OCT-2001; 200US-0296876P.
26-DEC-2001; 200US-033659P.
20-FEB-2002; 2002US-0338580P.
                                                                                                                                                                                                                                                                                                                                                                                                                                           characterizing gene function.
ADL99994 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                    26-MAR-2002; 2002WO-US009187.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                                                              Morrissey D, Maswiggen JA,
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MCSW/) MCSWIGGEN J A.
BEIG/) BEIGELMAN L.
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                                                                                                  Hepatitis B virus.
                                                                                                                     US2003206887-A1.
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                                    20-MAY-2004
                                                                                                                                      06-NOV-2003
                   ADL99994
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2002US-0408378P

Beigelman L;

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Virucide; Hepatotropic; Gene therapy; ss; short interfering nucleic acid; siNa; hepatitis B virus; HBV; RNA interference.
The antisense region 3'-terminal overhang is complementary to RNA encoding HBV. The siNA is useful for treating human hepatitis B virus infections, and for characterising pathways of gene function, e.g. to inhibit activity of target genes in a pathway to determine the function of uncharacterised genes in gene function analysis. The siNA molecules any also be used in clinical, industrial, environmental, agricultural and/or research settings. The present sequence represents 1 of 1504 HBV siNA molecules of the invention.
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                                                                                                                                                                                                                            0.4%; Score 14.4; DB 1; Length 19; 22.5%; Pred. No. 1.8e+03; ve 5; Mismatches 1; Indels
                                                                                                                                                                                    Sequence 19 BP; 4 A; 4 C; 4 G; 0 T; 7 U; 0 Other;
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ADM00600 standard; RNA; 19 BP.
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2002US-0363124P.
2002WO-US009187.
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2002US-0406784P.
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2000US-00636385.
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2001US-0296876P
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2001US-0337055P.
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                                                                                                                                                                                                                                                       62.5%;
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                                                                                                                                                                                                                                                       1 Similarity 62.5
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24-OCT-2000; 2
08-JUN-2001; 2
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24-OCT-2001;
05-DEC-2001;
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06-JUN-2002;
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(BEIG/) N
                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                           Virucide, Hepatotropic, Gene therapy, 8s, short interfering nucleic acid, siNA, hepatitis B virus, HBV; RNA interference.
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sina molecules of the invention.
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0.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 1.8e+03; ative 0; Mismatches 1; Indels Sequence 19 BP; 6 A; 3 C; 5 G; 0 T; 5 U; 0 Other; Query Match

Local Similarity 93.8 nes 15; Conservative

Best Loca Matches

ADL79493 standard; RNA; 19 ADL79493; RESULT 2000

20-MAY-2004 (first entry)

Human HER1 (EGFR) transcript target sequence/siNA upper strand, SEQ:658

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; dung screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; BGFR; HER1; c-erb-B-1; target sequence; ss.

Homo sapiens.

WO2003070912-A2

28-AUG-2003.

20-FEB-2003; 2003WO-US005045

11-MAR-2002; 2002US-0363124P. 29-MAY-2002; 2002WO-US016840. 06-JUN-2002; 2002US-00163552. 06-JUN-2002; 2002US-0386782P. 20-FEB-2002;

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regions, or alternatively are assembled from sense enube and antibense regions, or alternatively are assembled from a sense oligonucleotide. Specifically, the sinks include short interfering RNA (sirkNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (sirkNA). The sinks can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sink, conjugates and/or complexes of sink, and vectors that express sink. The sinks are corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The sinks are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, cumban might (EGFR) targeted double-stranded sink, which is identical to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HRR1, HRR2 HRR3 and HRR4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                        New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                 Jamison
                                                                                                                                                                                                                                                                                                                             Mcswiggen J, Pavco P, Beigelman L, Fosnaugh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 4 C; 7 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 658; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the HER1 transcript target sequence.
                       29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408778P.
09-SEP-2002; 2002US-0409233P.
19-SEP-2002; 2002US-00251117.
21-OCT-2002; 2002US-00251117.
   2002US-0393924P
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                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 75.0
Matches 12; Conservative
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The invention relates to a short interfering nucleic acid (siNA) molecule that down-regulates expression of a hepatitis B virus (HBV) gene by RNA interference or that inhibits HBV replication. Also disclosed are the following: (i) a method of modulating the expression of a HBV gene in a tissue explant; (ii)) a method of generating a library of siNA constructs having predetermined complexity; (iii) a cell containing one or more siNA molecules; (iv)a kit containing a siNA molecule which can be used to coganism; and (v) a method for synthesising a siNA molecule. The siNA molecule is adapted for use to treat HBV infection, and comprises a sense and an antisense region of a HBV target gene in a cell, tissue or complementary to an RNA sequence encoding HBV and the sense region comprises sequence complementary to the antisense region comprises the sense region and the sense region and the sense region and the sense region of the siNA molecule, where sense region and the comprises the sense region of the siNA molecule, where sense region and the sense region of the siNA molecule is a polynucleotide. Inher molecule is a polynucleotide inhibit or a innerse region comprise segion of the siNA molecule. The linker molecule is a polynucleotide. Inher molecule is a polynucleotide. Inher molecule is a polynucleotide. The antisense region comprises a 3'-terminal overhang and the antisense region comprises a 3'-terminal coverhang and the antisense region comprises a 3'-terminal coverhang and the antisense region comprises a 3'-terminal coverhang is conplementary to RNA cenceding HBV. The siNA is useful for treating pathways of gene function, e.g. to inhibit activity of traget genes in a pathway to determine the function coff uncharacterised genes in a pathway to determine the function and/or research settinings. The present sequence represents 1 of 1504 HBV sina molecules.
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Gaps ö 0.4%; Score 14.4; DB 1; Length 19; 5.0%; Pred. No. 1.8e+03; ve 3; Mismatches 1; Indels 1794 CCAGAGTGACGTCTGG 1809

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ઠ 셤 RESULT 2001

ADL79800 standard; RNA; 19 BP 20-MAY-2004 (first entry) ADL79800;

Human HER1 (EGFR) sinA lower strand, SEQ ID NO:965.

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER1; c-erb-B-1; ss. ADL 79800/C
XX
AC ADL 75
XX
XX
DT 20-M
XX
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XX
KW
KW
RNA

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03-JUN-2004 (first entry)
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                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAR-2004.
                                                                                                                                                                                                                                                31-JUL-2003,
                                                                                                                             ss; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor ceceptor (EGFR) genes (including HBR). HBR2 and HBR3 how HBR3 by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and matisense cregions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonuclectides, and can be chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Specifically synthesised. The invention also capacise to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of EGFR genes in cells, tissue explants or cyganisms (e.g., by ex vivo gene thorapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, such useful for trug screening, pharmacogenomics, contain upplying gene function, and gene mapping (e.g., of single nucleotide such and very or solve sinks are also expressed the sequence represents the lower strand of a breast and engage and gene mapping (e.g., of single nucleotide solve the such as breast and engage and gene mapping engages.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beigelman L, Fosnaugh K, Jamison S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 4 A; 7 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human HER1 (EGFR)-targeted double-stranded sina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 965; 171pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1794 CCAGAGTGACGTCTGG 1809
                                                                                                                                                                 2002US-0358580P.
2002US-0363124P.
2002WO-US016840.
2002US-00163552.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM77315 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                      19-SEP-2002; 2002US-00251117.
21-OCT-2002; 2002US-00277494.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                       2002US-0393924P.
                                                                                                                                                                                                                                                                                              29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                   09-SEP-2002; 2002US-0409293P.
                                                                                                                               20-FEB-2003; 2003WO-US005045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen J, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-697612/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                              WO2003070912-A2
          Homo sapiens.
                                                                                                                                                                                                                                   06-JUN-2002;
                                                                                                                                                                                                                                                                       03-JUL-2002;
                                                                                                                                                                       20-FEB-2002;
                                                                                                                                                                                                                29-MAY-2002;
                                                                                       28-AUG-2003
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셤 8

Query Match

Best Loca

ADM77315

RESULT 2002 ADM77315/c

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                                                                                  treating autosomal recessive polycystic kidney disease; PKHD1; DHPLC PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention comprises the amino acid and coding sequences of fibrocystin proteins. The DNA and protein sequences of the invention are useful for diagnosing and treating autosomal recessive polycystic kidney disease. The present DNA sequence represents a DHPLC FCR primer that was used to screen for mutations in the human fibrocystin (PKHDI) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P2V11 receptor; receptor; human; antianaemic; cardiovascular-gen.; CNS-gen;; respiratory-gen.; antiasthmatic; uropathic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated nucleic acid comprising a sequence encoding a fibrocystin polypeptide, useful for diagnosing and treating autosomal recessive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 1.8e+03; tive 0; Mismatches 1; Indels
Human fibrocystin (PKHD1) gene DHPLC PCR primer #23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 8 A; 9 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human purine P2Y11 receptor forward PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Torres VE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MAYO-) MAYO FOUND MEDICAL EDUCATION & RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; SEQ ID NO 70; 136pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Harris PC, Ward CJ, Rossetti S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2315 GICTGTGTGTGTGTGT 2330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM41325 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-SEP-2002; 2002EP-00020415.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-SEP-2002; 2002EP-00020415
                                                                                                                                                                                                                                                                                                                                                                                  23-JAN-2003; 2003WO-US002038
                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JAN-2002; 2002US-0351110P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FARB ) BAYER HEALTHCARE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptide, useful for di
polycystic kidney disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JUN-2004 (first entry)
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Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-618286/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                  human; fibrocystin;
                                                                                                                                                                                                                                                           WO2003062453-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CNS-gen.; respin
PCR; primer; ss.
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Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            RESULT 2005
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Matches
                                                                                                                                                                                                                                                                                                                                                                                      AAQ71320/
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8
                                                                                                                                                                                                                                                                                                                                     셤
                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is a forward primer for human P2Y11 receptor

(P2Y11R) ADM41309. The forward primer, a reverse primer ADM41326 and

probe ADM41327 were used in an example from the invention for expression

profiling in human haematopoietic bone marrow cells. Higher expression of

P2Y11R mRNA was observed in erythroid progenitor (CD71+) cells than in

early progenitor (CD34+) cells. Expression decreased during

marrow compartment. The invention is based on the finding of increased

expression of P2Y11R mRNA in erythroid cells. It provides methods of

cuing P3Y11R p01ypeptides to screen for agents useful in the treatment

and/or prophylaxis of anaemia, especially pure red cell aplasia, anaemia

of chronic renal failure, anaemia of endocrine disorders, congenital

dyserythropoietic anaemia, iron deficiency, congenital atransferrinaemia

confidentic pulmonary haemosiderosis (claimed), and for the treatment

of haematological and cardiovascular diseases, disorders of the

peripheral and central nervous system, chronic obstructive pulmonary

disorder (COPD), asthma, genito-urological disorders and inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                             Screening for therapeutic agents useful for the treatment and/or prophylaxis of, e.g. anemia, comprises detecting binding of test compound with polypeptide and selecting compounds which bind to the polypeptide as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying a subject at risk of breast cancer by detecting the presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       breast cancer; cytostatic; gene therapy; human; ss; primer; PCR; SNP;
single nucleotide polymorphism; NUMA1; FLJ20625; LOC220074;
chromosome 11q13.3; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Extend primer 2 used to genotype human NUMA1/FLJ20625/LOC220074 SNP.
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                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 14.4; DB 1; Length 19; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reneland R;
                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 1 A; 2 C; 10 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kammerer SM,
                                                                                                           Example 1; SEQ ID NO 17; 25pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Braun A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   3706 TGGTGGCCAGAGGTGT 3721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP46373 standard; DNA; 19 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                             93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 93.8
ses 15; Conservative
 Froehlen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SEQU-) SEQUENOM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nelson MR,
                       WPI; 2004-271670/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-441051/41
                                                                                    potential agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2004047623-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
 Kauschat D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                    diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roth RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 2004
                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                              The invention relates to a novel method for identifying a subject at risk
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                                                                                                                                                                                                                                    of breast cancer comprising detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a subject at risk of breast cancer, for early diagnosis, prevention and treatment of breast cancer, possibly via gene therapy, as well as to analyse and predict a response to a breast cancer treatment and in clinical drug trials. The current sequence is that of an Extend primer (also described as probe) the invention which was used to genotype human NUMAI/FLJ20625/LOC220074 region gDNA. FLJ20625 and LOC220074 have been mapped to chromosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ71320 was used as a primer in the PCR amplification of replication competent, recombined viruses (esp. retro- or adeno-viruses). This enabled said viruses to be detected in very small concentrations (eg. 1 in 100 000 cells), in body fulls and tissue samples (useful in gene therapy), and in viral vector preparations, cell lines and transduced target cells. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
of polymorphic variations in the ICAM, MAPK10, KIAA0861, NUMA1 or regions which are associated with breast cancer in a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting recombination of viral nucleic acid sequences - by recombined product with specific primers, esp. for detecting competent retro-viruses produced by gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Score 14.4; DB 1; Length 19; 33.8%; Pred. No. 1.8e+03; ved. No. insertes 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Recombinant viruses; retro-viruses; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 2 A; 8 C; 5 G; 4 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer for the detection of recombinant viruses.
                                                                                                                                            Example 7; Page 106; 289pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 37; 51pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2525 GGCAGGGAGCTGGGCC 2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ71320 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 GGCAGGGAACTGGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 93.8
nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         position 11q13.3
                                                                      from a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9419491-A1
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25-APR-1995
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Location/Qualifiers

/note= /*tag=

94WO-JP000725 93JP-00132739

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Amino group; immunoassy; antibody; detection; ss.
                                    Amino labelled oligonucleotide (3+).
                                                                                                                                                                                                                    (NISR ) NISSUI PHARM CO LID.
                 (first entry)
         (revised)
                                                                                                                                                                                                                                       Toyoda N;
                                                                                                     modified base
                                                                                                                                                                                                                                                                                                         to be assayed
                                                                                                                                                                                28-APR-1994;
                                                                                                                                                                                                  10-MAY-1993;
                                                                                                                                           WO9427150-A1
                                                                                                                                                             24-NOV-1994.
        25-MAR-2003
04-SEP-1995
                                                                          Synthetic
                                                                                                                                                                                                                                       Oku Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2008
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                                                                                                                                                                                             MDC protein; breast cancer; mamma carcinoma; ovary cancer; chromosome-17; primer; polymerase chain reaction; PCR_i ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RT-PCR was used to detect MDC expression in tissues of the human CNS (cerebrum, cerebellum and fetal brain) and in endocrine or reproductive organs (testis, ovary, mamma, adrenal gland, thymus and pancreas) using the primers given in AAQ76125 (bases 1764-1780 of the MDC cDNA sequence of AAQ76120) and AAQ76120 antisense, bases 1976-1957 of AAQ76120).
                                                                                                                                                                                                                                                                                                                                                                                                                        Novel MDC protein and DNA encoding it - used to develop prods. for the study, diagnosis and therapy of cancers, partic. breast and ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 0.4%; Score 14.4; DB 1; Length 20; Local Similarity 93.8%; Pred. No. 1.8e+03;
        0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; Page 18; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2097 CCAGGACACCCCCAGC 2112
                                               3781 ACACCTGGTTGCTAAC 3796
                                                                                                                                                                                                                                                                                                        93JP-00136602.
93JP-00257455.
94JP-00049904.
94JP-00073328.
                                                                                                                AAQ76126 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                        94EP-00107487.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 ccaedacaececcaec 16
                                                                                                                                                                                  Human MDC PCR primer BCO12
                                                                                                                                                               (first entry)
                                                             16 ACACCTGGTTGCTGAC
                            15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15; Conservative
                                                                                                                                                       (revised)
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                                                                                                                                                                                                                                                                                                                                                         (CANC-) CANCER INST
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-038478/06
                                                                                                                                                                                                                                                                                                                                                                                    Nakamura Y, Emi M;
                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                   22-SEP-1993;
23-FEB-1994;
12-APR-1994;
                                                                                                                                                     25-MAR-2003
01-AUG-1995
                                                                                                                                                                                                                                                                                        13-MAY-1994;
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                                                                                                                                                                                                                                                   EP633268-A2.
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                                                                                                                                                                                                                                Synthetic.
                                                                                                                                   AAQ76126;
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        Query Match
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                            Matches
                                                                                             RESULT 20
AAQ76126
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                                                        Simultaneous assay of several antigens or antibodies - using single immobilised nucleotide(s), and a mobile phase containing the complementary nucleotide(s) bound to a labelled ligand for the molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino labelled oligonucleotide 3(+).
                                                                                                                                                                                                                               Example 1; Page 16; 49pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3651 CTTGCTTGCCTGCAGG 3666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ75223 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 93.8
Matches 15; Conservative
WPI; 1995-006969/01.
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04-SEP-1995
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AAQ75221 standard; DNA; 20

RESULT 2007

AAQ7522

AAQ75221

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Primers T41001-T41382 are derived from novel human gene signature (GS) sequences which did not match with sequences deposited in Genbank release 76. The GS sequences (T19001-T26817) were obtained from 3.-directed cDNA libraries prepared from various human tissues; synthesis of CDNA was initiated from the 3.-end of mRNA by using poly(T) as the sole primer. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was determined (esp. using primers and probes derived from the GS sequences) as a means of diagnosing abnormal cell function or for recognising different cell types. The primers T41201-2 amplify clone pm1688 which comprises the GS HUMGS001467 (T20467), located on chromosome 19
                                                                                                                                                                                                                                                                                                                           Single-stranded DNA for identifying gene signatures - isolated from 3'-directed human cDNA library that reflects relative abundance of corresp. mRNA in specific human tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identification of pyrazinamide resistant mycobacteria - used to distinguish between M. bovis and M. tuberculosis; also methods for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; tive 0; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pncA gene; pyrazinamidase; PZAse; differentiation; treatmen
Mycobacterium bovis; Mycobacterium tuberculosis; infection;
PZA-resistant; probe; identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Probe for PZA-resistant Mycobacterium sp. pncA gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 5 A; 10 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Fig 8; 2245pp; Japanese.
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ID AAV03673 standard; DNA; 20 BP.
                                      93JP-00355504.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 93.0.
Thes 15; Conservative
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                                                                                                                                                                                                       Matsubara K, Okubo K;
                                                                                                     (MATS/) MATSUBARA K. (OKUB/) OKUBO K.
                                                                                                                                                                                                                                                                      WPI; 1995-206931/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-032662/03
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                                      12-NOV-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequences given in AAQ75219-25 are oligonucleotides which are used to demonstrate the method of the invention. These oligonuclectides may contain an amino group on their 5' end. The method constitutes an immunoassy of several immunochemical ligands. The reagent contains antibodies which specifically recognise one of the ligands to be assayed. Each antibody is attached to a different specific poly-or oligonucleotide, such as these. The antibodies are attached to a unitable label, e.g. horseradish peroxidase, biotin, avidin, digoxogenin, etc. The reagent reacts with the ligand to be assayed to give a complex which bears the label and also bears the nucleotide specific to the ligand. A complementary nucleotide is then added to which the complex becomes attached. The immobilised complex can then be assayed using the label. Several antibodies or antigens in a sample can be assayed using a single reagent. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Simultaneous assay of several antigens or antibodies - using single immobilised nucleotide(s), and a mobile phase containing the complementary nucleotide(s) bound to a labelled ligand for the molecule to be assayed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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Amino group; immunoassy; antibody; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
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nes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oku Y, Toyoda N;
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                                                                                                                                                                                                                                                                                                                                         10-MAY-1993;
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RESULT 2009

Query Match

Best Loca Matches

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Gaps

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treatment;

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Gaps

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0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels

93.8%;

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expression, e.g. inflammation. When labelled they can be used diagnostically to determine overexpression of ILIR, also to determine localisation and distribution of this expression for research, diagnostic
for treatment of disease in humans caused by excessive receptor
                                                                                                                                  Sequence 20 BP; 0 A; 11 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                              2587 GCGCTCGGCCCTCCC 2602
                                                                                                                                                                                                                                 Matches 15; Conservative
                                                                                              or therapeutic purposes
                                                                                                                                                                                                              Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ01480;
                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                           RESULT
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                                                                                   The present sequence is a probe for a pyrazinamide (PZA) resistant Mycobacterium sp. pncA gane, which encodes an altered pyrazinamidase (PZAse). A novel method for differentiating between M. bovis and M. tuberculosis comprises detecting in a sample an altered pncA gene encoding a His57Asp substitution indicating that the sample comprises M. bovis. The pncA gene and PZAse may be used to treat individuals infected with PZA-resistant M. tuberculosis. The pncA gene or PZase may be administered with at least one other antimicrobial agent, e.g. isoniazid or rifampicin. Probes corresponding to sections of the pncA gene may also be used to identify resistant strains or to distinguish between M. bovis and M. tuberculosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This is a novel oligomer comprising 20 covalently linked nucleotides which bind to the 5' untranslated region of the interleukin-1 receptor (ILIR) markh. Expression of ILIR, in cells and tissues can be modulated by compositions comprising oligomers which are able to specifically hybridise with target areas of its encoding sequence. The composition can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Type I interleukin-1 receptor; ILIR; human; IL1 protein; hybridisation; inflammation; ss; 5' untranslated region; phosphorothioate linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2'-substituted oligonucleotide(s) specific for interleukin-1 receptor type I - used to modulate expression and detect overexpression of the receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Phosphorothiate oligonucleotide sequence 8065 targeting ILIR mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/note= "Phosphorothioate internucleotide linkage"
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                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A; 7 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Geiger T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 19; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dean N,
           mycobacterial infection.
                                                        Claim 48; Page 50; 63pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  512 TGGAGCGCTCCCCCCA 527
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                                                                                                                                                                                                                                                                                                                                                                                                                                             93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV09181 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
nes 15; Conserv
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              treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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0.4%; Score 14.4; DB 1; Length 20;
Best Local Similarity 93.8%; Pred. No. 1.88+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer used to amplify an ORF of Chlamydia trachomatis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 1446; 1755pp; English.
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2 GCGCTCGGCTCCTCCC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-IB001939
                                                                                                                                                                                                           AAZ01480 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-371125/31.
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15; Conservative
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(PERM/) PERMUTT M A.
(INOU/) INOUE H.
                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                            WPI; 1999-181060/15.
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Matches 15; Conserv
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         Homo sapiens.
                           WO9909214-A1.
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                                                                                   13-AUG-1997;
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Synthetic
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                                                                                                                         Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                               New isolated baboon histo-blood group O allele - used to develop products for producing group O non-human primates for use in xenotransplantation of organs and tissues.
                                                                                                                                                                                                                                                                                                                                                                                                          AAV65179-V65183 are PCR primers used in the amplification of human ABO DNA which is used in a method for identifying baboon O alleles, and carriers of O alleles as well as methods for producing baboons and baboon cells, tissues and organs having a group O phenotype. Such cells, tissues and organs can be used for xenotransplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNKl; JNK2; Gell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
                                                                                                                                                     Blood group, O allele, baboon, carrier, phenotype, xenotransplantation, A allele, B allele, ABO, human, PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                             Szalay A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 10 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                             Fagoaga OR,
                                                                                                                                                                                                                                                                                         (UYLO-) UNIV LOMA LINDA MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JNK1-specific probe ISIS No: 12548.
                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 34; 50pp; English.
                                                                                                                                                                                                                                                                                                             Diamond D, Nehlsencannarella S,
                                                                                                                                  Human ABO DNA PCR primer Uaftx7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1977 GCCCTCCCAGAGGCCC 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX29317 standard; DNA; 20 BP.
         193 GAGGCTGAGGACACAG 208
                                                                          AAV65183 standard; DNA; 20 BP
                          1 GAGTCTGAGGACACAG 16
                                                                                                                                                                                                                                                    98WO-US009464
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                                                                                                               (first entry)
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                                                                                                               04-MAR-1999
                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                                  Synthetic
                                                                                              AAV65183;
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AAX29317
                                                      RESULT 2013
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                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides that detect and modulate the expression of un N-terminal kinase proteins - useful for treating hyperproliferative diseases and inhibiting tumor growth in animals, and for modulating protein protein phosphorylation by these proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer; WFS1; antidiabetic; renal; opthalmic; auditory; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to antisense oligonucleotides that detect and modulate the expression of Jun N terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding a JNK1, JNK2 or JNK3 protein, and which modulate expression of these proteins. The oligonucleotides are useful for modulating JNK protein. The oligonucleotides are useful for modulating JNK protein of a protein and eall cycle progression in cultured cells or animal cells. The oligonucleotides are phosphorylated by a JNK protein, and the expression of a cellular protein that promotes one or more metastatic expression of a cellular protein that promotes one or more metastatic events. The oligonucleotides also form pharmaccutical compositions for treating animals with a hyperproliferative disease, and for inhibiting tumor growth in an animal
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                                                                                                                                                                                                                                             Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 6 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                             Dean N, Monia BP, Nero PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 66; 190pp; English
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98WO-US016488.
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Murine leukemia virus.
    Permutt MA, Inoue H,
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          WPI; 2000-293106/25
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                                                                                                                                                                     01-JUL-1998;
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                                                                                                                                                           13-JAN-2000.
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                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA09033-34 are primers based on a predicted 5' untranslated region, based on the mouse EST sequence that was homologous to the human WFS1 cond based on the mouse EST sequence that was homologous to the human WFS1 cond based on the ware used for RT-PCR with MIN6 RNA. The WFS1 gene is from human forcomosome 4p, and is located between markers D4S50 and D4S431, conditions of which are associated with Wolfram syndrome. The nucleic acids may be used as a biological marker for early diagnosis of Wolfram CC acids may be used as a biological marker for early diagnosis of Wolfram CC syndrome and for predicting the predisposition of an individual to the syndrome. Disruptions of the gene which result in low protein expression or the expression of inactive protein products are associated with the development of the disease. The gene is also useful for gene replacement thereof agents (i.e. to rectify the mutations and/or supplement the individuals own production of the polypeptide) and for developing new methods and agents (i.e. agents which bind with either the nucleic acids or the protein to modulate it's expression and/or activity) for treating Wolfram syndrome. It is particularly useful for generating antibodies to the protein (i.e. Ab). Wolfram syndrome is a combination of familial juvenile conset diabetes mellitus, diabetes insipidus, optic atrophy and deafness
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                                                                                                                                        Nucleic acids associated with, and useful for the diagnosis and treatment of Wolfram syndrome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0
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site-specific integration; genomic rearrangement; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer used to amplify and mutate the MLV integrase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.4; DB 1;
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Klein D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BAVA-) BAVARIAN NORDIC RES INST AS
                                                 Mueckler M:
                                                                                                                                                                                                                           Example 1; Page 28; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       viral infection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            187 GAGGACGAGGCTGAGG 202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 gagaacgaggcrgagg 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ60549 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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es 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Salmons B,
(MUEC/) MUECKLER
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Control of the properties of the pol gene within the integrase region, was used in conjunction with PCR primers AA260550-58 (which are specific to the integrase region within the pol gene). The primers were used for deletion mutagenesis at the C-terminus of the primers were used for deletion mutagenesis at the C-terminus of the murine leukemia virus (MLV) integrase gene, beyond the catalytic site.

The resulting protein is inactivated. The mutated gene was used in the construction of an integrase deficient packaging cell line for the construction of an integrase deficient packaging cell line for the construction of an integrase deficient packaging cell line for the steroviral vectors of the invention. These retroviral vector comprise at site-specific integrate sequence that allows the site-specific integrate into a non-coding region of the host because they integrate specifically into a non-coding region of the host genome, rather than randomly, thus con-coding region of the host genome, rather than randomly, thus call minating the risk of causing deleterious genomic rearrangements. The derivoiral vectors (and systems containing them in packaging cells or derived retroviral particles (RVP)) are used for in vivo or in vitro gene therapy of viral infections. They are also used in the treatment of genetic, metabolic, proliferative or other diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense; gene therapy; JNK2 protein; apoptosis; cancer; cellular hyperproliferation; Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to antisense oligonucleotides (AAC62844-
C63000, AAA96093-A96099 and AAA07993) that hybridise specifically to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diabetes; Jun N-terminal kinase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 131; 160pp; English.
                            Example 3; Page 22; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3781 ACACCTGGTTGCTAAC 3796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC62860 standard; DNA; 20 BP.
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Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC62860;
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nucleotide encoding a Jun N-terminal kinase (JNK2) protein, resulting in decrease of JNK2 expression and leading to induction of apoptosis. The present sequence is one such antisense oligonucleotide. The oligonucleotides of the present invention are useful for treating diseases or conditions with reduced apoptosis, e.g. cancer and cellular hyperproliferation. The oligonucleotides may also be used to increase the stimulation of apoptotic proteins, e.g. for treating Alzheimer's or Parkinson's disease, amylotrophic lateral sclerosis, retinitis, pigmentosa, epilepsy, mycoardial infarction, stroke, obstructive jaundice, polycystic kidney and diabetes. The present sequence may have a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primers AAA61528-29 were used to amplify CDNA encoding a peanut resveratrol synthase polypeptide. The resveratrol synthase gene is used to produce edible transgenic plants which exhibit increased levels of resveratrol glucoside. Resveratrol is a stilbene, and a promoter of nitroxide formation, which is a vasodilatory agent and inhibits platelet aggregation. Transgenic plants of the invention are useful for increasing the nutritional value, for increasing disease resistance and for decreasing spoilage of the plant. Resveratrol synthase and resveratrol glucoside isolated from such plants are also useful for providing nutritional and pharmaceutical benefit to a human or animal by ingestion. Raised levels of resveratrol glucoside in foodstuffs such-as wines, may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New edible transgenic plant useful for nutritional supplements to inhibit cancer and cardiovascular diseases, is transformed with portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peanut; resveratrol synthase; edible plant; transgenic plant; resveratrol glucoside; stilbene; nitroxide formation; vasodilatory; platelet aggregation; nutritional value; disease resistance; spoilage; cardiovascular disease; cancer; artherosclerosis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for peanut resveratrol synthase polypeptide cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 6 C; 3 G; 6 T; 0 U; 0 Other;
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Best Local Similarity
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14-NOV-2000
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AAA63529,C
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DT 14-NOV-2
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be useful for the prevention of cardiovascular diseases, cancer, artherosclerosis and inflammation. (Updated on 15-5EP-2003 to standardise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to loblolly pine polynucleotides with one or more Simple Sequence Repeats (SSRs) (see AAA74205-A74322). SSRs are also known as microsatellite DNA repeats. The SSRs are useful as genetic markers for genetic mapping, population genetics studies and inheritance studies in various plant breading programmes. The present sequence is a PCR primer used for detecting the presence of a SSR locus in a pine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polynucleotide having simple sequence repeat useful as markers in plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for genetic characterization e.g. genetic mapping study, an inheritance study of a commercially important trait in a plant breeding program.
                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; loblolly pine; Simple Sequence Repeat; SSR; microsatellite DNA repeat; genetic marker; mapping; inheritance study; population genetics study; plant breeding programme; ss.
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                                                                                            Length 20
                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                         Reverse PCR primer for loblolly pine locus RIPPT914.
                                                           BP; 5 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                       0.4%; Score 14.4; DB 1;
33.8%; Pred. No. 1.8e+03;
ve 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 7 C; 3 G; 9 T; 0 U; 0 Other;
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                                                                                                       Best_Local Similarity 93.8%;
Matches 15; Conservative
                                                                                                                                                                           16 AGGTGCTGAATGGCTC 1
                                                                                                                                                                                                                                                               AAA74357 standard; DNA; 20
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                                                                                                                                                      59 AGGTGCTGAATGCCTC
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nes 15; Conservative
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(ECHT/) ECHT C S:
(NELS/) NELSON C D.
(USDA ) US SEC OF AGRIC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nelson CD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genomic DNA sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200042210-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-1999;
19-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pinus taeda.
                                                                                                                                                                                                                                                                                                                            29-NOV-2000
                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Si
Matches 15;
                              OS field)
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                                                                                                                                                                                                                                                                                            AAA74357;
                                                                                           Query Match
                                                                                                                                                                                                                               RESULT 2019
                                                                                                                                                                                                                                               AAA74357
 88888
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Human protein kinase C-theta; PKC-theta; PKCT; PRKCT; nPKC-theta; PRKCQ;

isozyme; serine/threonine protein kinase; signal transduction; calcium-independent function; JNK/SAPK pathway upstream activator; Jun N-terminal kinase/stress-activated protein kinase;
T-cell signalling pathway; cell cycle control; callular activation; API transcription factor activation; AIDS aetiology; apoptosis; cytoskeletal arrangement; proliferation; wound healing disorder; angiogenesis; insulin signalling; chromosome 10p15; angiotession inhibition; antisense; cancer; inflammation; diabetes; phosphorothioate; 2'-MOB gapmer; ss.

Homo sapiens.

US6190869-B1 20-FEB-2001.

Human protein kinase C-theta antisense oligonucleotide, SEQ ID NO:54.

04-MAY-2001 (first entry)

AAF59861;

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human insulin-like growth factor binding protein-5 (IGFBP-5). The invention relates to a composition for treatment of hormone-regulated cancer, comprising an antisense oligonucleotide (such as this sequence) which inhibits expression of IGFBP-5 by hormone-regulated tumour cells. The compositions is useful for delaying progression of hormone-regulated tumour cells such as prostatic cancer cells or breast cancer cells, to an androgen-independent state, by treating hormone sensitive tumour cells with the antisense sequence which inhibits expression of IGFBP-5 by the tumour cells. The composition can also be used for treating a hormone-cresponsive cancer in an individual, and administering the composition to the individual after initiation of hormone-withdrawal to induce apoptotic cell death of hormone-responsive cancer cells and therefore delaying the progression of hormone-responsive cancer cells to a hormone-independent state in the individual. It can also be used for inhibiting or delaying metastatic boney progression to inhibit the expression of IGFBP-5 by the hormone-responsive cancer cells to a hormone-independent composition to inhibit the expression of IGFBP-5 by the chormone-responsive cancer cells and therefore inhibiting or delaying metastatic boney progression of the tumour in a mammal, by administering the composition to inhibit the expression of IGFBP-5 by the concer cells, and therefore inhibiting or delaying metastatic boney progression of the tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Composition for treating hormone-regulated cancer, e.g. breast and prostatic tumors, comprising an antisense oligonucleotide that inhibits expression of insulin like growth factor binding protein-5 by hormone-regulated tumor cells.
                                                                                                                                                                                                                        Insulin-like growth factor binding protein-5; IGFBP-5; human;
antisense oligonucleotide; hormone-regulated cancer; prostatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; Ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure, Page 38, 45pp, English.
                                                                                                                                                                              Antisense IGFBP-5 inhibitor #31.
                                         AAA91225 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYBR-) UNIV BRITISH COLUMBIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUL-2000; 2000WO-CA000853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0144495P
                                                                                                                                                                                                                                                                       breast cancer; therapy; ss.
                                                                                                                                  08-MAY-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-168448/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MIYA/) MIYAKE H.
                                                                                                                                                                                                                                                                                                                                                             WO200105435-A2.
                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                         25-JAN-2001.
                                                                                   AAA91225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gleave M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
RESULT 2020
                       AAA9122
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Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human protein kinase C-theta useful for inhibiting expression of human protein kinase C-theta in human cells.

99US-00429322. 99US-00429322.

26-OCT-1999; 26-OCT-1999; Cowsert LM;

Bennett CF,

WPI; 2001-210378/21.

(ISIS-) ISIS PHARM INC.

Example 15; Col 43-44; 40pp; English.

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Sequences AAF59817-AAF59896 represent phosphorothioate 2'-MOE gapmer antisense targetted to the human protein kinase C-theta gene, which continated to the human protein kinase C-theta gene, which is the syrpession. The antisense oligonucleotides were designed to target different regions of the human protein kinase C-theta RNA, and were analysed for their effect on protein kinase C-theta RNA levels by committed two real-time PCR. Protein kinase C-theta RNA levels by CC theta, PKCT, PRKCT, nPKC-theta and PRKCQ) is one of several protein kinase C isozymes and is ubiquitously expressed, with the highest levels being found in haematopoietic cell lines. It has been shown to function in a calcium-independent fashion, and it is involved in a variety of contin a calcium-independent fashion, and it is involved in a variety of signal transduction pathways, for example, it is an upstream activator of the JNK/SAPK (UNIN N-terminal Kinase/Stress-activated protein kinase) compathway. Protein kinase C-theta is also involved in T-cell signalling compathways, cell cycle control, cellular activation, API transcription factor activation and the aetiology of AIDS, and has also been implicated and wound repair. It is additionally involved in insulin signalling and continue repair. It is additionally involved in insulin signalling and is thought to play a role in the development of diabetes in humans. The colligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with protein kinase C-theta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 0.4%; Score 14.4; DB 1; Length 20; Local Similarity 93.8%; Pred. No. 1.8e+03; es 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1908 CCGCATGGACAAGCCC 1923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 CCGCATGGACAATCCC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Gaps ö

2618 CCTGCAGGGAAGCCCC 2633

Best Local Similarity 93.8%;

15; Conservative

Matches

AAF59861 standard; DNA; 20 BP

RESULT 2021 AAF59861/C ID AAF5986

.; 0

MTR1; TRP-related protein; Ca2+ regulation; calcium regulation; tumor; transient receptor potential family; BWS; Beckwith-Wiedemann syndrome; 11pl5.5 abnormality; chromosome 11; anticancer; developmental activity; intracellular calcium ion regulation; hormone; growth factor; apoptosis; cell growth; cell death; cell differentiation; urogenital disease; polycystic kidney disease; calcium influx; Wilms tumor; rhabdoid tumor;

Location/Qualifiers

rhabdomyosarcoma; ds.

Homo sapiens.

'number= 19

intron

exon

ø

/number= 19

WO200132693-A2

10-MAY-2001

Д 11. .20 /*tag= *tag=

Prawitt D, Pelletier J, Zabel B;

WPI; 2001-316417/33.

(UYGU-) UNIV GUTENBERG JOHANNES.

99DE-01053167.

04-NOV-1999;

06-NOV-2000; 2000WO-DE003876.

Human MTR1 exon19/intron19 junction.

09-AUG-2001

AAH20613;

BP

AAH20613 standard, DNA; 20

RESULT 2023

AAH20613

20 AGGACGAGGATGAGGA 5

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Sequences AAG92560-C92639 represent antisense oligonucleotides targetted to the human nucleolin gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human nucleolin mRNA, and were analysed for their effect on nucleolin mRNA concleolin mRNA, and were analysed for their effect on nucleolin mRNA levels by quantitative real-time PCR. Nucleolin (also known as P92 or C23) is the most abundant nucleolar phosphoprotein in actively growing cells. Nucleolin primarily participates in ribosome biogenesis and transport of ribosomal components, being able to transiently bind to preciposomes in the nucleolus via a ribonucleoprotein consensus sequence. However, it has also been shown to be involved in cytokinesis, on ucleogenesis, cell proliferation and growth, transcriptional represence to a member of the Ag-NOR (active ribosomal genes and whose expression is associated with the prediction cyganiser region) family of proteins which are markers of active ribosomal genes, and whose expression is associated with the prediction of tumour growth rate. The presence of antibodies against nucleolin are associated with systemic connective tissue diseases such as systemic connective diseases. The oligonucleotides of the invention are useful for diagnosis, connective diseases. The oligonucleotides of the invention are useful for diagnosis, connective diseases of the invention are useful for diagnosis,
                                                                                                                                    Human nucleolin phosphorothioate antisense oligonucleotide, SEQ ID NO:31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound targeted to human nucleolin which specifically hybridizes with and inhibits the expression of human nucleolin, useful for modulating the expression of nucleolin in cells.
                                                                                                                                                                            Human nucleolin, P92; C23; phosphoprotein; ribosome biogenesis; ribosome transport; cytokinesis; nucleogenesis; cell proliferation; cell growth; transcriptional repression; replication, signal transduction; chromatin decondensation; Ag-NOR family; nucleolin antibody; systemic connective tissue disease; SLE; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                   scleroderma-like chronic graft versus host disease;
expression inhibition; tumour formation; cancer; inflammation;
immune disorder; phosphorothioate; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; Col 41-42; 41pp; English
AAC92581 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-00433699.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-00433699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment of
                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bennett CF, Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-079848/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prevention and
                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                         27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       JS6165786-A
                                            AAC92581;
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This invention describes a novel DNA sequence (I) encoding the WTR1

protein that: (i) has at least one biological activity of a TRP

(cransient receptor potential) family protein; (ii) is connected with

etiology of BWS (Beckwith-Wiedemann syndrome) and/or (iii) is connected

with tumors involving lip15. Saboramalities. The products of the

cripulation of intracellular calcium ion levels, which are essential for

cellular responses to hormones and/or growth factors; also in apoptosis

and cell growth, death and differentiation, and in urogenital diseases,

including polycystic kidney disease. (I) and related ribozymes, antisense

CR RNA, proteins and antibodies (Ab) are used to treat or prevent diseases

CR RNA, proteins and antibodies (Ab) are used to treat or prevent

CR protein, or with calcium influx into cells, e.g. BWS, Wilms tumor,

rhabdoid tumors and rhabdomyosarcoma. Probes from (I), or Ab, are also

used for diagnosis of such diseases. (I) can also be used for recombinant

production of MRT1 proteins (II) (used for analysis, characterization and

cherapy), as tissue or chromosomal markers, for identifying genetic

chardy antibodies. (II) are used to raise anti-protein or

chardy Antibodies. (II) as tissue markers; for identifying

competitive assays for (II), as tissue markers; for identifying

competitive assays for (II), as tissue markers; for identifying

chreapents human MTRI gene exon 19/intron 19 junction region described in DNA encoding MTR1 protein, useful e.g. for treating Beckwith-Wiedemann syndrome and tumors, also related proteins and antibodies. Disclosure; Fig 9; 46pp; German. ö expression, such as tumour formation, immune disorders and inflammation Gaps with nucleolin ; 0 Query Match 0.4%; Score 14.4; DB 1; Length 20; Best Local Similarity 93.8%; Pred. No. 1.8e+03; Matches 15; Conservative 0; Mismatches 1; Indels conditions associated Sequence 20 BP; 2 A; 9 C; 1 G; 8 T; 0 U; 0 Other;

188 AGGACGAGGCTGAGGA 203

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Gaps

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Indels

Length 20;

Query Match

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Best Loca Matches

à g ABK92361;

RESULT 2024

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The present invention relates to a method of detecting tumour cells, by detecting extranuclear DNA consisting of a single-strand with a double-stranded stem-loop structure containing at least 2 binding sites for DNA binding proteins. The method can be used to detect cancer cells in tissue sections, biopsies, body fluids etc. and can be used in the diagnosis and monitoring of cancer. The present sequence is an extranuclear DNA sequence capable of being detected using the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting tumor cells from presence of specific stem-loop DNA molecules outside the nucleus, useful for diagnosis and monitoring of tumors.
                                                                                                                                                                                                                                                                                                                                                                         extranuclear DNA; stem-loop; DNA-binding protein; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bound_moiety= "nucleotides 15-12 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12. .15
/*tag= f
/bound_moiety= "nucleotides 7-4 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= b
/bound_moiety= "nucleotide 17 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= g
/bound_moiety= "nucleotide 2 of itself"
0.4%; Score 14.4; DB 1;
33.8%; Pred. No. 1.8e+03;
ve 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                      Cancer cells detection oligonucleotide M29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "DeltaEF1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'bound_moiety= "CHOP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 21; Page 13; 27pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bound moiety=
                                                                               1188 GCTGACCCTGGGCAAG 1203
                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-SEP-2000; 2000DE-01046318.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-SEP-2000; 2000DE-01046318.
                                                                                                      93.8%;
                                                                                                                                                                                                                      AAL45762 standard; DNA; 20
                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                  tumour cell detection; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ಹ
                      Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Abken H, Schinkoethe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . .20
*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..10
*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-331116/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ABKE/) ABKEN H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DE10046318-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                           Нощо варіепв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein_bind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein_bind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           misc binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     misc binding
                                                                                                                                                                                                                                                                                                    27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     stem_loop
                                                                                                                                                                                                                                                               AAL45762;
                                                                                                                        0
    Query Match
                                                                                                                                                                                                                                                                                                                                                                                 Cancer;
                                                                                                                                                                                   RESULT 2025
                                                                                                                                                                                                        AAL45762,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
                                                                                                                                                                                                                                          엄
                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a novel CaCo 2 cell line showing an enhanced expression of CYP3A4 without the addition of an inducer for CYP3A4 corpusation of an inducer for CYP3A4 is the major form of cytochrome P450 expressed in the capression. CYP3A4 is the major form of cytochrome P450 expressed in the colline also comprises a method for growing this cell line and a culture medium. The cell line of the invention may be used as a rool in pharmacoctor of the intestinal barrier and may be used as a tool in pharmacoctor toxicology. The cell line is also useful in in vitro models for screening the bioavailability of pharmaceutical compounds, xenobiotics or contronmental pollutants. The invention also discloses RT-PCR primers that may be used to analyse the expression of genes with sequence similarity to CYP2A4 such as CYP2A7, CYP3A4, CYP3A5, NADPH reductase, cyP2C, CYP2B1, CYP2B6, CYP2B1, CYP1A1, MDR1, MDR3, MRP1, MRP2, MRP3, GST CYP2C, CYP2B1, CYP2B7, UGT1A6 and beta actin in the cell line of the invention of one of the above mentioned containing the cell line of the invention under different culture
                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CaCo 2 cell line having enhanced expression of CYP3A4 without the addition of an inducer for CYP3A4 expression, useful in in vitro models for screening the bloavailability of pharmaceutical compounds, xenobiotics or environmental pollutants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer; reverse transcription; RT-PCR; ss; human; CaCO2; CYP1A1; MRP1; CYP3A4; intestinal barrier; pharmaco-toxicology; xenobiotic; MRP3; environmental pollutant; CYP3A7; CYP3A4; CYP3A5; NADPH reductase; MRP2; CYP2C; CYP2B1; CYP2D6; CYP2E1; MDR1; MDR3; GST Pi; GST alpha 4; UGT2B7; UGT1A6; beta actin.
                                                                                                                            Gaps
                                                                                                                            ö
                                                                                     Length 20;
                                                                                                                          Indels
                                          Sequence 20 BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                   Score 14.4; DB 1;
Pred. No. 1.8e+03;
                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 12; 38pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYLO-) UNIV CATHOLIQUE LOUVAIN.
                                                                                                                                                                 2731 GGGTACCTGAAGATGG 2746
                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                            Human CYP1A1 RT-PCR primer #2.
                                                                                     Match 0.4%;
Local Similarity 93.8%;
Les 15; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-MAY-2000; 2000EP-00870108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           L7-MAY-2000; 2000EP-00870108
                                                                                                                                                                                                        16 gegraccreaacaree 1
          the method of the invention
                                                                                                                                                                                                                                                                                                          ABK92361 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Burteau N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-407056/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP1158045-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schneider Y,
                                                                                                                                                                                                                                                                                                                                                                                     16-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-NOV-2001.
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Sequence 20 BP; 6 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

conditions

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The sequences given in ABA00075 and ABA00083-85 are probes and primers which were used in the detection of H. pylori infection pre- and post-
contractment in a patient, compared to the presence of human DNA. These sequences may be used in the method of the invention for detecting a H. pylori infection. The method comprises: (a) determining an integrity of a Helicobacter pylori nucleic acid present; or (b) amplifying and detecting a felicobacter pylori nucleic acid at least 200, at first, second or third Helicobacter pylori nucleic acid at least 200, infection if the integrity of the nucleic acid exceeds a predetermined infection if the integrity of the nucleic acid exceeds a predetermined contracter pylori infection, determining its status, monitoring mucleic acids are detected. The method is useful for detecting a Helicobacter pylori infection, determining its status, monitoring disease by detecting a high-integrity Helicobacter pylori nucleic acid in a pasays detecting a high-integrity Helicobacter pylori uncleic acid in a pasays detecting H. pylori infection usually lack the specificity to distinguish a successfully treated patient from a patient with a continuing H. pylori infection. The present invention of non-invasive method uses more sensitive and more specific assays to test for and to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting, grading and/or monitoring a Helicobacter pylori infection by detecting a high-integrity H. pylori nucleic acid in a patient sample.
                                                                                                                                   Primer; probe; detection; Helicobacter pylori; integrity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Page 27; 28pp; English.
                                                                                                                                                                                                                                                                                                                               05-JAN-2001; 2001US-00755004.
                                                                                                                                                                                                                                                                                         04-JAN-2002; 2002WO-US000267.
                                                         25-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                     EXAC-) EXACT SCI CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-599807/64.
                                                                                             Human APC primer #1
                                                                                                                                                                                                             WO200259379-A2
                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                   01-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                         Shuber AP;
                     ABA00084;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention comprises antisense oligonucleotides that are capable of inhibiting expression of the human clusterin gene. The antisense oligonucleotides of the invention are useful for inhibiting the expression of clusterin in cells. The antisense oligonucleotides are also useful for treating an animal with a disease or condition associated with clusterin (e.g., hypercholesterolaemia; cardiovascular disorders; hyperproliferative disorders; and hyperlipidemic disorders; has present DNA sequence represents a clusterin antisense oligonucleotide of the invention. NOTE: The present DNA sequence has a phosphorothioate backbone and also contains 2'-O-methoxyethyl wings
                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel antisense compound targeted to nucleic acid molecule encoding clusterin, useful for treating animal having disease associated with clusterin such as hyperlipidemic disorder, cardiovascular disorder.
                                                                                                                                                                                                                                                                                                                                                                                                       Human, antisense inhibition, antisense oligonucleotide, clusterin; hypercholesterolaemia, cardiovascular disorder; ss; hyperproliferative disorder; hyperlipidemic disorder; phosphorothioate, backbone; 2'-0-methoxyethyl wing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; Ive 0; Mismatches 1; Indels
                                   Ouery Match
0.4%; Score 14.4; DB 1; Length 20;
Best Local Similarity 93.8%; Pred. No. 1.8e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                     Human clusterin inhibiting antisense oligonucleotide 52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 4 A; 8 C; 6 G; 2 T; 0 U; 0 Other;
Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 84; 125pp; English.
                                                                                                                 3651 CTTGCTTGCCTGCAGG 3666
                                                                                                                                                                                                                                                   ABN99718 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-SEP-2001; 2001WO-US028235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-SEP-2000; 2000US-00659791
                                                                                                                                                   17 crrecarecerecaes 2
                                                                                                                                                                                                                                                                                                                               16-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Freier SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-404805/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200222635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monia BP,
                                                                                                                                                                                                                                                                                           ABN99718;
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                                                                                                                                                                                                                                                                                                                                                                                      Xenobiotic response element; XRE; transcription factor; drug screening;
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0
                                                               0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; tive 0; Mismatches 1; Indels
monitor H. pylori infection and course of treatment
                                  Sequence 20 BP; 5 A; 12 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     Xenobiotic response element PCR primer.
                                                                                                                                                                                                                                                      ABN83781 standard; DNA; 20 BP
                                                                                                                                    856 GAGGAGCTGGTGGAGG 871
                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                    15; Conservative
                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                        02-SEP-2002
                                                                                                                                                                                                                                                                                       ABN83781;
                                                                     Query Match
                                                                                                                                                                                                                                       ABN83781
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Gaps

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2407 CTGGGTGTCCCCCCTG 2422

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20

93.8%;

Local Similarity 93.8 es 15; Conservative

Best Loca Matches

ABA00084 standard; DNA; 20 BP

RESULT 2027 ABA00084/c ID ABA0008

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The invention provides a novel system for evaluating transcription activity in eukaryotes using novel reporter vectors designated as intranscriptional regulation analysis system (TRAS). In an example from the invention, a TRAS-XRE vector was constructed comprising a reporter gene, a minimal promoter and a S-repeated xenobiotic response element (XRE, see PRN813777), and was used to transfect HeLa and HepG2 cell lines. The present sequence is a primer that was used in a PCR to evaluate establishment of TRAS-XRE cell lines. Such cell lines can be used to examine the activity of transcription factors and to determine the effects of substances affecting transcription activity. These compounds can be used to treat or prevent a disease or disorder caused by dioxinlike compounds, such as endocrine disruption, decrease of sperm number,
                                                                                                                                                                                                                                                                                                                                                            Screening substances affecting transcription associated with promoter, by measuring protein expressed by a cell transfected with a reporter vector comprising reporter gene encoding protein and the promoter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        decrease of testis weight, endometriosis, breast cancer, testis cancer, prostate cancer, sexual organ malformation, decrease of immune function
dioxin; cytostatic; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 4; Page 41; 103pp; English
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                                                                                                                                                            06-NOV-2001; 2001WO-KR001881
                                                                                                                                                                                                    06-NOV-2000; 2000KR-00065656
                                                                                                                                                                                                                                                                                 Chang M, Shin H,
                                                                                                                                                                                                                                                                                                                        WPI; 2002-500130/53.
                                                                                                                                                                                                                                         (MEDE-) MEDEXBIO CO
                                                                             WO200237077-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and hypospadia
                                     Unidentified
                                                                                                                    LO-MAY-2002
                                                                                                                                                                                                                                                                                 Lee S,
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; 0 0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; Ive 0; Mismatches 1; Indels Sequence 20 BP; 4 A; 3 C; 6 G; 7 T; 0 U; 0 Other; 1007 TGCACAGATCTCCCG 1022 1 Similarity 93.8%; 15; Conservative 16 TTCACAAGATCTCCCG 1 Query Match Best Local Similarity Matches 셤

; 0

Gaps

ACC44846 standard; DNA; 20 04-JUN-2003 (first entry) ACC44846;

BP.

Human antibody 146B7 VH region PCR primer #2.

Human; antibody; interleukin 15; IL-15; IL-15 receptor; antirheumatic; antiarthritic; antiinflammatory; antipsoriatic; immunosuppressive; cytostatic; antimicrobial; psoriasis; arthritis; rheumatoid arthritis; inflammatory bowel disease; cancer; transplant rejection; infectious disease; 146B7; PCR primer; ss.

Synthetic

WO2003017935-A2

06-MAR-2003

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The present invention describes an isolated human monoclonal antibody (I) that specifically binds to human interleukin 15 (IL-15) and inhibits IL-15 to the specifically binds to human interleukin 15 (IL-15) and inhibits IL-15 induced proinflammatory effects upon binding to its receptor. (I) has antirheumatic, antiarthritic, antiinflammatory, antipsoriatic, immunosuppressive, cytostatic and antimicrobial activities. (I) is useful for inhibiting IL-15 induced, but not IL-2 induced, TNFalpha production by T cells or monocytes, which involves contacting IL-15 induced, real or proliferation, which involves contacting IL-15 induced, T cell or contacting IL-15 induced, but not IL-2 induced, T cell or such inhibiting IL-15 induced, but not IL-2 induced, T cell or cells such as peripheral blood mononuclear cells (I) in the presence of T cells such as peripheral blood mononuclear cells (PBMCs) or CTLL-2 cells. (I) is useful for treating or preventing a disorder mediated by carbitish, inflammatory bowel disease, cancer, transplant rejection or an infectious disease. (I) is also useful for diagnosing an IL-15 can infectious disease. (I) is also useful for diagnosing an IL-15 cell expressing IL-15 by contacting the presence of IL-15 antigen, or a complex between antibody or its portion and IL-15 and a control sample, complex, where formation of a complex, where formation of a complex is indicative of the presence of IL-15 in the sample. The present sequence represents a PCR complex for the human antibody 14687 VH region, which is used in an
                                                                                                                                                                                                                                                                                                        New isolated human monoclonal antibody that specifically binds to human interleukin-15 and inhibits IL-15 induced proinflammatory effects, useful for treating or preventing rheumatoid arthritis and psoriasis.
                                                                                                                                                                               Gerritsen AF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 3 C; 8 G; 4 T; 0 U; 2 Other;
                                                                                                                                                                            Van Dijk MA, Schuurman J,
                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Page 62; 114pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               example from the present invention
                        23-AUG-2002; 2002WO-US026769.
                                                                        23-AUG-2001; 2001US-0314731P.
                                                                                                                                                                                                                                                          WPI; 2003-312792/30.
                                                                                                                                                                            Van De Winkel JGJ,
                                                                                                                           (GENM-) GENMAB INC
                                                                                                                                                                                                       Baadsgaard O;
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Gaps ö 0.4%; Score 14.4; DB 1; Length 20; 75.0%; Pred. No. 1.8e+03; 3; Indels 2; Mismatches 15; Conservative Local Similarity Query Match Matches

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Telomere; cancer; age; telomerase; male infertility; primer; ss. Primer XpYp-415AT used for determining telomere length. (first entry) 22-APR-2003

ВР

ABZ68355 standard; DNA; 20

RESULT 2030

ABZ68355

ABZ68355;

WO2003000927-A2. Homo sapiens.

21-JUN-2002; 2002WO-GB002855. 03-JAN-2003

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Gaps

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Indels

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21-SEP-2001; 2001GB-00022755.
                                                                                                                                                                                                                              20-JUN-2002; 2002WO-US019516
                                                                                                                                                                                                                                      20-JUN-2001; 2001US-0299664P
                                                                                                     diagnosing male infertility
                                                                                                                                                               ABZ68414 standard; DNA;
                                                                                                                                                                                                                                                      Carpten J, Trent JM,
                          WEI; 2003-175294/17.
                                                                                                                        Local Similarity
                                                                                                                                                                                                               WO2003000112-A2.
                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                               22-APR-2003
                                              infertility.
                                                                                                                                                                                                                      03-JAN-2003
                                                                                                                                                                       ABZ68414;
                                                                                                                     Query Match
                                                                                                                                                        RESULT 2031
                   Baird
                                                                                                                             Matches
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PCR primers ABZ68114-15 were used to amplify a fragment of the human Rhase L gene. The amplified fragment was examined for the presence of a mutation, in the course of the invention. The specification describes a method for detecting cancer or a predisposition to cancer in a mammal. The method comprises detecting at least one mutation in a gene encoding interferon inducible 2',5'-oligoadenylate-dependent Rhase L in a test sample obtained from the mammal, where at least one mutation indicates cancer or predisposition to cancer. The method is useful for producing a medicament for treating a mammal prophylactically or therapeutically for cancer that is due to a complete or partial loss of wild-type RNase L 1..20
/*tag= a
/mod base= OTHER
/mote= "This oligonucleotide has a phosphorothioate
/note= "This oligonucleotide has a thosphorothioate
backbone and 2-'methyoxyethyl (2.'MOE) wings at the 5'
and 3' ends, which are 5 nucleotides in length. Also all
cytidine residues are 5-methylcytidines" yase gene expression, useful for preventing or treating a developmental lisorder or aberrant apoptosis, e.g. infection, inflammation or tumor antisense oligonucleotides for modulating sphingosine-1-phosphate Cytostatic, antimicrobial; antiinflammatory; tumour; infection; sphingosine-1-phosphate lyase; developmental disorder; apoptosis; inflammation; antisense; phosphorothioate; ss. Sphingosine-1-phosphate lyase antisense oligonucleotide, SEQ ID 0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other; 0; Mismatches cocation/Qualifiers 1308 BP 26-SEP-2002; 2002WO-US030575. 28-SEP-2001; 2001US-00967669. cereaagerecreaaa 17 ACC70595 standard; DNA; 20 (first entry) 1293 CGTGAAGATGCTGAAA Conservative (ISIS-) ISIS PHARM INC. WPI; 2003-381581/36. Local Similarity Les 15; Conserv WO2003028637-A2 Key modified_base 13-AUG-2003 10-APR-2003 Bennett FC, formation. Synthetic. Query Match ACC70595; disorder RESULT 2032 Matches ************ ò 임 ö Determining telomere length of mammalian chromosomal DNA by annealing and covalently binding telorettes, useful in assessing a potential anticancer treatment and/or assessing, treating or diagnosing male The present sequence represents a primer, which is used in the method of the invention for determining telomere length. The method comprises annealing the 1 end of a single-stranded oligonucleotide to a single-stranded overhang comprising TIAGG repeat sequences, and covalently binding the oligonucleotide to the ? end of the C-rich telomeric strand forming a ligation product. This ligation product is amplified to form a primer extension product, whose length is then detected. The method is useful in assessing potential anti-cancer treatments and cancer-related procedures. It can also be used in assessing telomere dynamics with age, effect of modulating telomerase activity, and in assessing, treating or Gaps Stephan ô PCR primer used to amplify a 461 bp fragment of RNase L gene. Length 20; 1; Indels Isaacs WC, Human; RNase L gene; cancer; mutation; PCR; primer; Sequence 20 BP; 4 A; 5 C; 5 G; 6 T; 0 U; 0 Other; Score 14.4; DB 1; Pred. No. 1.8e+03; 0; Mismatches 1; Smith J, Walsh PC, (USSH) US DEPT HEALTH & HUMAN SERVICES (UYWA-) UNIV WALES COLLEGE OF MEDICINE. Claim 23; Page 36; 50pp; English. 3782 CACCTGGTTGCTAACC 3797 ВР 23-JUN-2001; 2001GB-00015451 0.4%; 16 caccreerrearacc 1 (first entry) 15; Conservative WPI; 2003-175260/17.

The present invention relates to novel antisense oligonucleotides (ACCYOS20-ACCYOS27) which are targeted to a sphingosine-1-phosphate lyase DNA sequence, and specifically hybridizes with the nucleic acid and inhibits the expression of sphingosine-1-phosphate lyase. The antisense

Example 15; Page 74; 118pp; English.

Detecting cancer or predisposition to cancer in a mammal for producing a medicament for treating a mammal for cancer that is due to a complete or partial loss of wild-type RNase L by detecting at least one mutation in

Example 2; Page 19; 32pp; English

RNase

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31-JUL-2002; 2002WO-US024370.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotides which hybridizes to, and modulates the expression of Jun N-terminal kinase, useful for treating a disease or condition characterized by a reduction in apoptosis, e.g. prostate cancer, inflammation or fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an oligonucleotide (antisense, AS) comprising 8-30 nucleotides connected by covalent linkages, where the oligonucleotide has a sequence specifically hybridisable with a nucleic acid encoding a Jun N-terminal kinase (JNK) protein and modulates the expression of the JNK protein. Also included are a pharmaceutical composition comprising
      condition associated with sphingosine-1-phosphate lyase, particularly a developmental disorder, or a disease or condition arising from aberrant apoptosis, e.g. infection, inflammation or tumour formation
                                                                                                                                                                                                                                                    Human Jun N-terminal kinase, JNK1, antisense oligonucleotide ISIS12548.
oligonucleotides are useful for treating an animal having a disease or
                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                  conjunctival scarring, hyperproliferative disease; cancer; probe
                                                                                            ;
0
                                                                                                                                                                                                                                                                         88; human; Jun N-terminal kinase; JNK1; JNK2; JNK3; antisense;
                                                                      0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                   cytostatic; antiinflammatory; apoptosis; prostate cancer; prostate tumour; inflammation; fibrosis; fibrotic disease; fibrotic scarring; peritoneal adhesion; lung fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                   /note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaarde WA
                                                  Sequence 20 BP; 8 A; 4 C; 1 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                 3007 TIGITITAAAACIGGA 3022
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98US-00130616.
99US-00287796.
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                                                                                                                                                                                        ADA26564 standard; DNA; 20
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                                                                                 Best Local Similarity 93.8
Matches 15; Conservative
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(DEAN/) DEAN N M.
(MONI/) MONIA B P.
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modified_base
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                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-APR-1999;
15-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-AUG-1998;
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                                                                                                                                                                                                             ADA26564;
                                                                        Query Match
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(NERO/)
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the AS oligonucleotide (or its bioequivalent, and a pharmaceutical carrier), treating an animal having/suspected of having/prone to having a hyperproliferative disease (by administering to a prophylactic or therapeutic amount of the composition of the AS oligonucleotide).

C therapeutic amount of the composition of the AS oligonucleotide, modulating the expression of a JNK protein in cells or tissues by contacting the cells or tissues with the AS oligonucleotide, modulating the cell cycle progression (or the phosphorylation of a protein properson or more metastatic events in cultured cells, inhibiting common or animal) by administering the oligonucleotide to the cells, inhibiting an animal) by administering the oligonucleotide to the cells, inhibiting the growth of a tumour in an animal by administering a human having a disease or condition associated with a JNK protein or characterised by a reduction in apoptosis by administering a prophylactic or therapeutic amount of the condition characterised by a reduction in apoptosis, such as a disease or condition characterised by a reduction in apoptosis, such as a disease or condition (e.g. fibrotic scarring, peritoneal adhesions, lung fibrosis or condition (e.g. fibrotic scarring, peritoneal adhesions, lung fibrosis or condition (e.g. fibrotic scarring, peritoneal adhesions, lung fibrosis or condition, such as cancer. The antisense oligonucleotides may also be used as research agents and diagnostic aids, to detect the present of converse of sources of sources
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REI silencing transcription factor; COREST; antisense oligonucleotide;
developmental; hyperproliferative; disorder; neuronal cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       study the function of one or more genes in the animal. The present sequence is an antisense oligonucleotide targeting human JNK1.
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/note= "all cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indel8
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/note= "2'-O-methoxyethyls (2'MOE)
16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a new antisense compound comprising 8-50 nucleobases in length targeted to a nucleic acid molecule encoding a crepressor for REI silencing transcription factor (CoREST), where the compound specifically hybridises with and inhibits the expression of COREST. The COREST antisense oligonucleotide has any of 72 specifically claimed sequences of 20 bp, given in the specification. The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of COREST, such as a developmental disorder and/or a hyperproliferative condition like neuronal cancer. The current sequence represents an antisense oligonoleotide for the
                                                                                                                                                                                                                                                                                               New antisense oligonucleotide compounds, useful for the diagnosis, prevention and/or treatment of conditions with aberrant expression or activity of COREST, such as developmental and/or hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer, male infertility; infertility-associated mutation, azoospermia factor; Y-chromosome; cystic fibrosis transmembrane conductance regulator; CTFR; Kallmann syndrome; KALI, androgen resistance; steroid 21-hydroxylase; CYP21; microarray; quantitative trait locus; in vitro fertilization;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; SEQ ID NO 14; 145pp; English
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                               01-AUG-2001; 2001US-00920671.
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Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                   Bennett CF, Freier SM;
                                                                                                   (ISIS-) ISIS PHARM INC.
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This invention describes a novel method for investigating genetic infertility or predisposition in males. The method involves selecting at least two infertility-associated mutations which are recessive or least two infertility-associated mutations with the heterozygous state and/or only in the homozygous state. Preferably at least one azoospermia factor is detected which may be lost by microdaletions in intervals 5 or 60 the Y-chromosome. Also any of several hundred mutations, listed, present in the cystic fibrosis transmembrane conductance regulator (CTFR), Kallmann syndrome (KALI), androgen resistance (AR) or steroid 21-hydroxylase (CYP21) genes may be detected. Probes for the mutated genes and/or native nucleic acid, or their complementary strands, are fixed to a carrier, particularly as a microarray, then tested for hybridization with oligonucleotides from or synthesized from, a patient sample and hybridization detected. Multifactorial analysis is by standard statistical methods, particularly the quantitative trait locus method. The method is used to diagnose the quantitative trait locus method. The method is used to diagnose the quantitative trait locus method. The method is used to diagnose the quantitative for possible application of the subjects with oligospermia for possible application of the intracytoplasmic sperm injection method. Analysis of many mutations improves diagnosis of the genetic basis of male infertility, in cluding complex interactions between different heterozygotic mutations. A chip for analyzing genetic infertility in males comprises oligonucleotides that represent known mutations (nonsense or missence, insertions, and propermy of the method in the propermy of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fibrosis transmembrane conductance regulator, Kallmann syndrome, androgen resistance and steroid 21-hydroxylase genes. ADD42140-ADD42633 represent oligonucleotides used in the microarray described in the method of the invention. NOTE: there are no SEQ ID's 133, 472 or 473 represented in the
                     Investigating male genetic infertility, useful for diagnosis e.g. for assessing suitabilty for in vitro fertilization, based on multifactorial analysis of infertility-related mutations.
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5-aminolevulinic acid synthase; enzyme; haem; P450 cytochrome; chicken;
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                                                                                                                            Claim 13; SEQ ID NO 407; 110pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID list of the specification
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Best Local Similarity
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Bertram

Tanzi RE,

Wang X, Ta Blacker DL;

Elliott KJ, Sampson AJ,

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Neurodegenerative disease; uPA; SNCG; IDE; KNSL1; LIPA; TNFRSF(Alzheimer's disease; neuroprotective; nootropic; gene therapy; Chromosome 10; PCR; primer; ss.
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08-NOV-2001; 2001US-0338010P.
09-NOV-2001; 2001US-033835P.
24-DEC-2001; 2001US-0337652P.
28-MAR-2002; 2002US-0368919P.
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(GEHO ) GEN HOSPITAL CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003054143-A2
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                                                                                       Meyer UA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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AC ADE4:
ADE43758/C
AC ADE4:
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The present invention relates to a method (M1) for determining a predisposition for or the occurrence of neurodegenerative disease in a saughject. The method comprises detecting in a target nucleic acid obtained from the subject the presence or absence of an allalic variant of one or more polymorphic regions of one or more genes selected from upA (Urokinase plasminogen activator). SNGG (gamma-synuclein), IDE (insulingerase), and TNFRSF6 (Tumour Necrosis Factor Receptor-SF6), where the presence of at least one of the allalic variant of one or more polymorphic regions is indicative of a predisposition for or the occurrence of neurodegenerative disease. The genes are all located on chromosome 10. M1 is useful for determining a predisposition for or the occurrence of, and for treating neurodegenerative disease, particularly Alzheimer's disease. The present sequence is a PCR primer, which was used
                                                                                                                                                                               Determining a predisposition for or the occurrence of neurodegenerative disease, e.g. Alzheimer's disease by detecting in a target nucleic acid the presence or absence of an allelic variant of one or more polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                 Example 3; Page 300; 848pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in the method of the invention.
                          Velicelebi G,
Mullin KM,
                                                                                                                       WPI; 2003-559131/52.
                                                           Saunders AJ,
                                 Becker KD,
                                                                                                                                                                                                                                                                                         regions.
The present invention describes an isolated nucleic acid sequence comprising at least a DR-4 nuclear receptor binding site, where the nucleic acid sequence functions as transcriptional enhancer of the 5-caminolevulinic acid synthase gene. Also described: (1) a genetic construct comprising the above nucleic acid sequence operably linked to a nucleic acid encoding a reporter molecule; and (2) a method for testing compounds for medulation of haem and/or P450 cytochrome synthesis, computating sultable cells comprising the above genetic construct with a test compound and detecting enhanced or repressed expression and/or transcription of the nucleic acid sequence encoding the reporter gene. The nucleic acid or genetic construct can be used for testing of chemical compounds as modulators of haem and/or P450 cytochrome synthesis. The present sequence represents a PCR primer used in the generation of a probe for the chicken 5-aminolevulinic acid synthase (AlaAS1) gene, which is used in the exemplification of the
                                                                                                                                                                                                                New transcriptional enhancer of the 5-aminolevulinic acid synthase gene, useful for testing of chemical compounds as modulators of heme and/or P450 cytochrome synthesis, comprises at least a DR-4 nuclear receptor
                                                                                               Zumsteg A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 4 C; 10 G; 1 T; 0 U; 0 Other;
                                                                                            Podvinec M,
                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 31; 69pp; English.
                                                                                            Kaufmann MR,
                                                                                            Fraser DJ,
                                                                                                                                                           WPI; 2003-877033/81
                                 (UYBA-) UNIV BASEL
                                                                                                                                                                                                                                                                                                                   binding site.
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                                                                                            Gaps
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Query Match 0.4%; Score 14.4; DB 1; Length 20; Best Local Similarity 93.8%; Pred. No. 1.8e+03; Matches 15; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kaufmann MR, Podvinec M,
                                                                                                                                                                                        2214 ACAATGTGAGGGGTCC 2229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF18169 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-2002; 2002WO-IB001258.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-2002; 2002WO-IB001258.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ALAS1 gene; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                          19 ACAATGTGAGGGCTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fraser DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-877032/81.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF18169;
                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 2038
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The present sequence is that of a forward PCR primer for the chicken aminolevulinic acid synthase (ALAS1) gene. The primer was used to generate a probe for ALAS1, in a PCR using chicken embryo liver genomic DNA as template. ALAS1 drug responsive enhancer sequences (ADRES) have been identified ADP18139-ADP18140. These include DR4 and DR5 nuclear receptor binding sites. An isolated nucleic acid sequence, such as the ADRES sequence, that comprises at least a DR4 site is claimed. A claimed genetic construct comprises the nucleic acid sequence and a reporter modulation of heam and/or cytochrome P450 synthesis. Cells compounds for modulation of heam and/or cytochrome P450 synthesis. Cells comprising the genetic construct are contacted with the test compound, and enhanced/reduced expression and/or transcription of the reporter gene is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Variant detecting primer extension assay extension primer, SEQ ID No 35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               variant detection; primer extension assay; mutation; cancer;
heterogeneous; sporadic mutation; genotyping; pooled sample; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting a variant in a primer extension assay, useful for analyzing
                                                                                                                                                                                                                                                                                            Score 14.4; DB 1; Length 20;
Pred. No. 1.8e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 4 C; 10 G; 1 T; 0 U; 0 Other;
            Disclosure; SEQ ID NO 31; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Whitney D;
                                                                                                                                                                                                                                                                                                                                                           2145 CCACGACCTGCTGCCC 2160
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                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 93.8%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-697649/66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF53079;
                                                                                                                                                                                                                                     detected
                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 2039
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primer extension assay, useful for analysing molecular events for identifying mutations indicative of cancer, by contacting a target nucleic acid primer complementary to a region of the target nucleic acid betecting a variant in a primer extension assay comprises contacting a target nucleic acid primer complementary to a region of the target nucleic acid, and extending the primer in the presence of a first uncleded that is complementary to a first variant nucleotide suspected to be at a position downstream of the region and a second nucleotide that

The invention relates to a novel method for detecting a variant in a

Example 5; SEQ ID NO 35; 54pp; English.

molecular events for identifying mutations indicative of cancer, by contacting a target nucleic acid primer complementary to a region of the target nucleic acid.

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is complementary to a second variant nucleotide at the position, thus to reduce misincorporation of the first nucleotide on a template comprising the second variant nucleotide. The methods are useful for analysing molecular events for identifying individuals with mutations indicative of cancer. They are particularly useful in detecting a rare mutation in a heterogeneous biological sample (e.g. sporadic mutation in a heterogeneous patient sample), detecting rare genotypes in genotyping reactions (e.g. viral genotyping reactions), or detecting mutant or viral sequences in pooled samples (e.g. detecting polymorphisms or inherited sequence variations in pooled patient samples). This polymucleotide sequence represents a primer used as part of the primer extension assay
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ansal airway dysfunction and a second active agent comprising an antiinflammatory steroid and bujquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antianthatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense; lung dysfunction; nasal airway dysfunction;
                                                                                                                                                                                                                                                                                                       Score 14.4; DB 1; Length 20;
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 12 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Li Y, Sandrasagra A, Ke
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВР
                                                                                                                                                                                                                                                                                                                                                                                        856 GAGGAGCTGGTGGAGG 871
                                                                                                                                                                                                                                                                                                           0.4%;
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                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                                                                                                                                             the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ85455;
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immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, antisense; lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antianfmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                   1; Indels
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                                                                                                                                                                                                      Sequence 20 BP; 10 A; 3 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                    0.4%; Score 14.4; DB 1; 33.8%; Pred. No. 1.8e+03;
                                                                                                                                                                         at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PDE4C oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 i Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                   3258 AAGATATTTTATTTGC 3273
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                                                                                                                                                                                                                                                                                                                               16 AAGTTATTTTATTTGC 1.
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                                                                                                                                                                                                                                                      93.8%;
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                                                                                                                                                                                                                                                   Local Similarity 93.8
les 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ99173;
                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                            RESULT 2041
                                                                                                                                                                                                                                                                   Matches
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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

Disclosure; SEQ ID NO 14415; 872pp; English.

ubiquinone

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, ? or 3' end genomic flanking regions, s' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or masal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,

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immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at tp. whoo.int/bub/Dublished or seminance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                                                                                                                                                                                                                     0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03;
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Tang L, Shahabuddin S;
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Homo sapiens. 17-0CT-2003 31-OCT-2002. Miller S, ABZ89955; Query Match 3 RESULT 2043 Matches Nyce ABZ89955/ 1D ABZ85/ AXX ABZ85/ XXX ABZ85/ XXX ABZ85/ DDE Human EWW anttiin EWW anttiin EWW aden EWW aden EWW 1009 XX Homo XX Homo XX Homo XX EPIC XX EPIC XX EPIC XX EPIC XX WPI; XX WPI; XX WPI; XX COTTING END OF COTTING XX COTTING EVEN OF COTTING XX COTTING EVEN OF COT 8866666666668888 ò 셤

immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO WPI; 2003-229219/22. WO200285308-A2. Homo sapiens. 17-0CT-2003 ubiquinone. Miller S, ABZ86363; Query Match Nyce JW, RESULT 2044 Matches ABZ86363/ ŝ 셤 ö lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences Pharmaceutical composition for treating ailments associated with impaired immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid Gaps Aguilar D; ö 0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; 1; Indels Pabalan J, Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other; 0; Mismatches Disclosure; SEQ ID NO 5197; 872pp; English. Katz B, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Human oligonucleotide sequence. 2633 CACATGTCCAGCACCT 2648 BP. 1 cácarcricaccarcr 16 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P (EPIG-) EPIGENESIS PHARM INC ABZ89955 standard; DNA; 20 (first entry) 15; Conservative WPI; 2003-229219/22. Local Similarity ri Y, WO200285308-A2 ubiquinone

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                                                                                                                                                                                                                                                                                                                                                                                                                       Human, antisense, lung dysfunction, nasal airway dysfunction; antisinflammatory steroid; ubiquinone; antisinflammatory; antiallergic; antisathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                      Length 20;
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                                    Sequence 20 BP; 1 A; 4 C; 9 G; 6 T; 0 U; 0 Other;
                                                                    0.4%; Score 14.4; DB 1;
93.8%; Pred. No. 1.8e+03;
tive 0; Mismatches 1;
at ftp.wipo.int/pub/published_pct_sequences
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Tang L, Shahabuddin S;
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The invention relates to a novel pharmaceutical composition, which has a

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,

immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data dor this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or Human, antisense; lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; Gaps Katz E, Pabalan J, Aguilar D; ő 0.4%; Score 14.4; DB 1; Length 20; 13.8%; Pred. No. 1.8e+03; 1; Indels Seguence 20 BP; 4 A; 8 C; 6 G; 2 T; 0 U; 0 Other; 0; Mismatches lung inflammation; respiratory disease; ds. Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Human oligonucleotide sequence. 1394 ACCTGCTGGGCGCCTG 1409 BP. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC. 93.8%; 19 Accrecrececacine 4 ABZ85454 standard; DNA; 20 (first entry) Local Similarity 93.8 nes 15; Conservative WPI; 2003-229219/22 WO200285308-A2. Homo sapiens. 17-OCT-2003 31-OCT-2002 ubiquinone Miller S, ABZ85454; Query Match Nyce JW, Matches à 유

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antieense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ansal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,

Claim 15; SEQ ID NO 696; 872pp; English.

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ö immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition, Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5 or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ansal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, Human; antisense; lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Gaps Katz E, Pabalan J, Aguilar D; ö . Match 0.4%; Score 14.4; DB 1; Length 20; Local Similarity 93.8%; Pred. No. 1.8e+03; les 15; Conservative 0; Mismatches 1; Indels Sequence 20 BP; 14 A; 2 C; 1 G; 3 T; 0 U; 0 Other; at ftp.wipo.int/pub/published_pct_sequences Disclosure, SEQ ID NO 6976; 872pp; English. Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Human oligonucleotide sequence. 3275 ABZ91734 standard; DNA; 20 BP 24-APR-2001; 2001US-0286137P. 23-APR-2002; 2002WO-US013135. (EPIG-) EPIGENESIS PHARM INC. w (first entry) 20 GTTATTTTATTTGCTT 3260 GATATTTTATTTGCTT WPI; 2003-229219/22. WO200285308-A2. Homo sapiens. 17-OCT-2003 31-OCT-2002. ubiquinone. Miller S, ABZ91734; Query Match Nyce JW, RESULT 2046 Matches ABZ91734/c ò g

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immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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Sequence 20 BP; 12 A; 0 C; 1 G; 7 T; 0 U; 0 Other;

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Human oligonucleotide sequence.
    ABZ86378 standard; DNA; 20
            (first entry)
            17-OCT-2003
        ABZ86378;
RESULT 2047
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Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 1620; 872pp; English.

which has a The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,

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preventing a respiratory, lung or malignant disease or condition, also preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject s tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
immunosuppressive, and cytostatic activity. The composition may have a
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Sequence 20 BP; 5 A; 6 C; 9 G; 0 T; 0 U; 0 Other;

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Best Local Similarity 93.89
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674 5 GCAGCAAGGAGGGCCC 20 659 GCAGCAAGGTGGGCCC Š g

ABZ88135 standard; DNA; 20 BP. ABZ88135; RESULT 2048 4BZ88135

(first entry) 17-0CT-2003

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, d Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Willer S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ansal airway dysfunction and a second active agent comprising an antinflammatory steroid and bujquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,

WO200285308-A2.

Disclosure; SEQ ID NO 3377; 872pp; English.

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immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory diseases or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published pct sequences
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Sequence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;

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ABZ91731 standard; DNA; 20 BP.

Human oligonucleotide sequence. (first entry) 17-0CT-2003 ABZ91731; RESULT 2049
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Human, antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antisathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Katz E, Pabalan J, Shahabuddin S; Sandrasagra A, Li Y, Sal Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 6973; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 5 and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ansal altrway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,

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use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject stissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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RESULT 2050 AAD47315,

AAD47315 standard; DNA; 20

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AAD47315;

(first entry) 24-FEB-2003 Human RT-PCR reverse primer for synaptophysin DNA isolation.

Human; insulin-secreting cell; neurogenin 3; ngn3; precursor stem cell; pancreatic exocrine cell; transplantation; RT-PCR; primer; ss.

Homo

WO200274946-A2.

26-SEP-2002

26-FEB-2002; 2002WO-DK000130.

26-FEB-2001; 2001US-0271474P.

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ö Gradwohl Serup P, Heimberg H,

WPI; 2003-018804/01.

Generating insulin-secreting cells from precursor stem cells or adult panoreatic exocrine cells, for generating glucose sensitive insulin secreting beta cells for transplantation, comprises using neurogenin3 or NeuroD/beta2

Example 5B; Page 36; 66pp; English.

The invention relates to a method for generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells. The method acid molecule encoding neurogenin 3 (ngm3) or exocrine cells to: a nucleic acid molecule encoding neurogenin 3 (ngm3) or NeuroD/Deta2; or an activator of ngm3 or NeuroD/Deta2 gene expression, under conditions effective to generate the insulin-generating cells from the precursor or exocrine cells. The invention is useful in generating insulin-secreting cells from precursor stem cells or adult pancreatic exocrine cells is useful for generating glucose sensitive insulin secreting beta cells suitable for transplantation, and for in situ development of insulin-secreting premature differentiation of precursor stem cells into insulin-secreting beta cells and for indentifying compounds that prevent or activate beta

ВЪ.

ABD22608 standard; DNA; 20

RESULT 2052

(first entry)

29-JUL-2004

ABD22608;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel isolated protein (1) comprises the amino acid sequence of wild type cryopyrin of 1034 amino acids, with the proviso that amino acid 198 is not Val, amino acid 352 is not Ala, amino acid 434 is not Ala, amino acid 627 is not Glu, or amino acid 703 is not Gln. The methods are useful for determining the presence of a disorder, treating inflammation, Familial cold uriticaria/familial cold autoinflammatory syndrome (FCV)FCAS) or Muckle Wells Syndrome (WWS), and identifying a substance useful in modulating binding of a cryopyrin protein to a second protein. The oligonuclocide is useful in diagnosing a disorder characterized by an aberrant CLASI gene. This sequence corresponds to a primer used to amplify the cryopyrin gene of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated cryopyrin protein and encoding nucleic acid, useful for diagnosing and treating inflammatory disorders, in particular familial cold urticaria, familial cold autoinflammatory syndrome and/or Muckle
present sequence is human RT-PCR primer for
                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         primer; antiinflammatory; cryopyrin; inflammation;
llial cold urticaria; familial cold autoinflammatory syndrome;
                                                                                                                     ö
                                                                                  0.4%; Score 14.4; DB 1; Length 20;
93.8%; Pred. No. 1.8e+03;
                                                                                                                 1; Indels
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                                                  Sequence 20 BP; 7 A; 6 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                          Human cryopyrin cDNA sequence primer #18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 18; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (LUDW-) LUDWIG INST CANCER RES
                 isolation of synaptophysin DNA
                                                                                                                                                   1281 TGTCACCGTAGCCGTG 1296
                                                                                                                                                                                                                                                                        ADM34346 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-OCT-2002; 2002WO-US031502
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                                                                                                  93.8%;
                                                                                                                                                                                    0
                                                                                                                                                                           17 TGTCACCGTGGCCGTG
                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                     15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hoffman H, Kolodner R;
cell differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Muckle Wells Syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-393448/37.
                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003031639-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Familial cold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                         ADM34346;
                                                                                    Query Match
                                                                                                                                                                                                                                   Matches
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comprising oligonuclectides, a novel composition, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and cantification or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition and/or bonofoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated in the lungs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchiis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system
                                                                                                                                                                                                        respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cyetic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel composition (a) a first active agent,
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                                                                                                                                                                                      Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted tnucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pabalan J, Aguilar D;
                                                                                                                                         Human cathepsin C-derived oligo SEQ ID 1620.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; SEQ ID NO 1620; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ii Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bronchodilating agent.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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Gaps

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Score 14.4; DB 1; Length 20; Pred. No. 1.8e+03; 0; Mismatches 1; Indels

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15; Conservative

Matches

0.4%;

Query Match Best Local Similarity

2318

2303 CACAGAGCTTTGGTCT

CACAGAGCTGTGTCT 17

셤 8

inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respirator distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system

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or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung

Thu Oct 28

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (B) the oligonucleotides, (C) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a caradrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered to reduce the production composition composition
e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted inucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pabalan J, Aguilar D;
                                                                                                                                      ö
                                                                                                 Length 20
                                                                                                                                      1; Indels
                                                        Sequence 20 BP; 5 A; 6 C; 9 G; 0 T; 0 U; 0 Other;
                                                                                               Score 14.4; DB 1;
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                    R49144-derived oligonucleotide SEQ ID 5197.
                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 5197; 763pp; English
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                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                            659 GCAGCAAGGTGGGCCC 674
                                                                                               0.4%;
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                                                                                                                                                                                                                                                                                                                ABD26185 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                               Query Match
Best Local Similarity 93.8
Matches 15; Conservative
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Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptor
                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide containing less percentage of adenosine, targeted unucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                               °,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense
                                                                                                                                                                                                                                                             0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human myosin X-derived oligonucleotide SEQ ID 1933.
                                                                                                                                                                                                                                   Sequence 20 BP; 1 A; 4 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A, Katz E,
J, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; SEQ ID NO 1933; 763pp; English.
                                                                                                                                                                                                                                                                                                                              2103 CACCCCCAGCTCCAGC 2118
                                                                                                                                                                                                                                                                                                                                                                                                                                          ABD22921 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-APR-2001; 2001US-0286036P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                  20 CAGCCCCAGCTCCAGC
                                                                                                                                                                                                                                                                                                 15; Conservative
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Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                 Best Local Similarity
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABD22921;
                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 2054
                                                                                                                                                                                                                                                                                                 Matches
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888888888888888888888888
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The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenargic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, circliammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system cere, into the system cere, the oligomucleotides into products that free adenosine into the system cere of the oligomucleotides the oit.
surfactant depletion or hyposecretion, when administered to a mammal. The objognucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.
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Sequence 20 BP; 4 A; 8 C; 3 G; 5 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; 1; Indels 0; Mismatches 93.8%; Query Match Best Local Similarity 93.8 Matches 15; Conservative

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2633 CACATGTCCAGCACCT 2648 1 CACATGTCCAGCATCT 16 g

ABD24365 standard; DNA; 20 BP 29-JUL-2004 ABD24365; RESULT 2055

(first entry)

AI672565-derived oligonucleotide SEQ ID 3377.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

Aguilar D;

WPI; 2003-093058/08

The invention describes a novel composition of a livite active agent, comprising oligomucleotides, effective for alleviating active agent, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and craduating adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligomucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) analgesic, hypotensive, immunosuppressive and cytostatic activity, is a certain a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production and/or bornchoconstriction and/or lung corresponding to inflammation, allergies and/or bronchoconstriction and/or lung with a disease or condition such as pulmonary viscometricition, allergies as sthma, impeded respiratory disease, pulmonary corresponding to inflammation, allergies asthma, impeded respiratory diseases, pulmonary corresponding to transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymication beautometrian and thereby, to prevent the programmation and thereby, etc, tissue environment and thereby, to prevent any unwanted effects due to it composition (a) a first active agent, Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted tnucleic acids associated with lung airway or lung dysfunction, and Claim 15; SEQ ID NO 3377; 763pp; English. invention describes a novel bronchodilating agent.

Seguence 20 BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;

Gaps ö 0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; 1; Indels 0; Mismatches 93.8%; Matches 15; Conservative Best Local Similarity Query Match

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ABD21684 standard; DNA; 20 BP ABD21684; RESULT 2056 ABD21684/c

Human stanniocalcin-derived oligo SEQ ID 696.

29-JUL-2004 (first entry)

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery of the invention also describes a kit, that comprises: (a) a delivery is a constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, consposition cancerastic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchost transcribed and an antial and an a
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                                                                                                                                                                                                                                                                    Katz E, Pabalan J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 696; 763pp; English.
                                                                                                                                                                                                                                                                                                  Shahabuddin S;
                                                                                                                                                                                                                                                             Li Y, Sandrasagra A,
Tang L, Shahabuddin
                                                                 23-APR-2002; 2002WO-US013143.
                                                                                                                           24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                               EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bronchodilating agent.
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31-OCT-2002
                                                                                                                                                                                                                                                             Nyce JW, ]
Miller S,
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ô Gaps ö 0.4%; Score 14.4; DB 1; Length 20; 13.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels 3260 GATATTTTATTTGCTT 3275 93.8%; 15; Conservative Local Similarity Query Match Matches

ABD32204 standard; DNA; 20 BP. Ŋ 20 Griaritiariacii RESULT 2057 ABD3220 셤

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(first entry) 29-JUL-2004 ABD32204;

Human PDE4C-derived oligonucleotide SEQ ID 14415.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cautedrant depletion or hyposecretion, when administered to a mammal. The colligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention an antiallersic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction are associated inflammation, allergies and/or surfactant hypoproduction are associated
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                    surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycotatic; cystic fibrosis; beta-adrenagic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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respiratory tract inflammation; adenosine sensitivity; lung; cancer;
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nucleic acids associated with lung airway or lung dysfunction, and
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Best Local Similarity 93.8%; Fred. No. 1.869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; SEQ ID NO 14415; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nyce JW, Li Y, Sandrasagra A, Ka
Miller S, Tang L, Shahabuddin S;
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating adenosine sensitivity, levels of adenosine (A) or (A) receptors, reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction also describes a kit, that comprises; (a) a delivery of the invention also describes a kit, that comprises; (a) a delivery of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytosetal cartivity, is a composition comparises oligonise. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligoner content of the anti-sense oligoner content for the anti-sense oligoner content of the anti-sense oligoner content
                                                                                                                                                                                                                                       Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                                                           AA497002-derived oligonucleotide SEQ ID 6973.
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Tang L, Shahabuddin S;
                                              ABD27961 standard; DNA; 20 BP
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                                                                                                                                             (first entry)
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                                                                                               ABD27961;
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RESULT 20
ABD27961/
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comprising oligonacieotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung of dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory districtses syndrome; allergic rinititis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                                                           Score 14.4; DB 1; Length 20;
Pred. No. 1.8e+03;
0; Mismatches 1; Indels
                                           Sequence 20 BP; 10 A; 0 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human stanniocalcin-derived oligo SEQ ID 697.
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prevent any unwanted effects due to it
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i, Shahabuddin S;
                                                                                                                                                                             2829 TACATATATATATA 2844
                                                                                                                                                                                                                                                                                                                                 ВЪ.
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                                                                                                           93.8%;
                                                                                                                                                                                                                      19 TTCATATATATATA 4
                                                                                                                                                                                                                                                                                                                                 ABD21685 standard; DNA; 20
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                                                                               Query Match
Best Local Similarity 93.8
Matches 15; Conservative
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Tang L,
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Miller S,
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pulmonary observation, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic thinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary the reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc. tissue environment and thereby, to prevent any unwanted effects due to it
amount of target polypeptide present in the lungs. The
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0.4%; Score 14.4; DB 1; Length 20; ilarity 93.8%; Pred. No. 1.8e+03; Conservative 0; Mismatches 1; Indels 3258 AAGATATTTTATTTGC 3273 Local Similarity es 15; Conserv Query Match Best Loca Matches ð

Gaps

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16 AAGTTATTTTATTTGC 1

ABD22593 standard; DNA; 20 BP. (first entry) 29-JUL-2004 ABD22593;

Human cathepsin C-derived oligo SEQ ID 1605.

Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; emphysema; chronic obstructive puimonary uisease pulmonary transplantation rejection; ss; primer.

Homo sapiens

WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Miller S, Nyce JW,

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense obligonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 1605; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchecometriction, respiratory tract inflammation, allergies and reducing adenosine ensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The

WPI; 2003-093058/08

Configuration of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The instructions are also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) adelivery confirmation has antiallergic, antialflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to collmonary obstruction, and/or bronchocomstriction and/or lungs. The pulmonary obstruction, and/or bronchocomstriction and/or lungs of inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, continued and allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary contransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligon corresponding to thymidines present in the target RNA serves to prevent the breakdown of the plantarion into predict that free adenosine into preducts that free adenosine into the present in the target RNA serves to prevent the breakdown of the present in the present should be anti-stone and the present and p e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it Gaps oligonucleotides are derived from a gene encoding or regulating ö 0.4%; Score 14.4; DB 1; Length 20; 13.8%; Pred. No. 1.8e+03; .ve 0; Mismatches 1; Indels Seguence 20 BP; 4 A; 8 C; 6 G; 2 T; 0 U; 0 Other; AA497002-derived oligonucleotide SEQ ID 6976. 1394 ACCTGCTGGGCGCCTG 1409 BP 93.8%; ABD27964 standard; DNA; 20 (first entry) 19 Accrecredecaccre 15; Conservative Local Similarity 29-JUL-2004 ABD27964; Query Match Matches ABD27964/ ð

Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Aguilar D; Pabalan J, pulmonary transplantation rejection; ss; primer. Katz E, Nyce JW, Li Y, Sandrasagra A, Miller S, Tang L, Shahabuddin (EPIG-) EPIGENESIS PHARM INC. 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. WO200285309-A2. Homo sapiens. 31-OCT-2002.

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This invention describes a novel composition, all title active agent, comprising oligomuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and creducing adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligomuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or treating a respiratory, lung or malignant disease. The administered correduction and/or bronchoconstriction and/or lung or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung or inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and or many and or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and or surfactant any unmentent of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine of the oligonucleotides into expect due to it tissue environment and thereby, to the prevent any unwanted effects due to i
                                                                                                                                                                                                                          invention describes a novel composition (a) a first active agent,
                             oligonucleotide containing less percentage of adenosine, targeted unucleic acids associated with lung airway or lung dysfunction, and
Pharmaceutical composition for treating asthma, has antisense
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                                                                                                                                                    Claim 15; SEQ ID NO 6976; 763pp; English
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Gaps ö Score 14.4; DB 1; Length 20; Pred. No. 1.8e+03; 0; Mismatches 1; Indels 2829 TACATATATATATA 2844 93.8%; Query Match
Best Local Similarity 93.0.
These 15; Conservative ઠ

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25-MAR-2004 (first entry)

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ADG27988 standard; DNA; 20 BP 11-MAR-2004 (first entry) ADG27988; RESULT 2062

Rice variaty-related PCR primer SeqID103.

rice variety; rice genome; quality control; adulterant detection; small trader protection; consumers protection; PCR; primer; ss.

Oryza sativa

18-DEC-2003

WO2003104491-A1

10-JUN-2003; 2003WO-JP007332

10-JUN-2002; 2002JP-00168875

(PLAN-) PLANT GENOME CENT CO LTD.

Determining a predisposition for or the occurrence of neurodegenerative

Tanzi RE

Ellliott KJ, Wang X, Mullin KM, Sampson AJ;

Velicelebi G,

Becker KD, Bertram L,

Saunders AJ,

WPI; 2004-060538/06.

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which comprises judging base types at any of 28 sites selected from the rice genome, or of the bases in these sites of base pairs of the bases at such sites in the complementary strand and relating the judged base types with the rice variety. The method is useful in distinguishing and with the rice variety. The method is useful in distinguishing and specifying varieties analogous to each other at DNA level for quality control of 24 rice varieties, to detect adulterants and mixing with cheaper varieties for protection of small traders and consumers.
                                                                                                                                                                                                                                           This invention relates to a novel method of distinguishing rice varieties
                                                                                                                     Distinguishing rice varieties based on polymorphism sites as markers to show different patterns in combination, useful in distinguishing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                        Nemoto H, Ideta O;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                   Disclosure, SEQ ID NO 103; 193pp; Japanese.
                                                                                                                                                              specifying varieties for quality control.
                                        Suzuki J, Ohta R,
(NAAG-) NAT AGRIC RES ORG JAPAN.
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Best Local Similarity
                                        Monna L,
                                                                             WPI; 2004-053624/05
                                        Minobe Y,
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human; neurodegenerative disease; urokinase plasminogen activator; uPA; gamma-symuclein; SNC3; insulin degrading enzyme; IDE; kinesin-like protein 1; KNSL1; lysosomal acid lipase; LIPA; tumour necrosis factor receptor SF6; TNFRSF6; Alzheimer's disease; PCR; Human neurodegenerative disease-related sequencing primer SeqID363 2001US-0336929P. 2001US-0338010P. 2001US-0338363P. 2001US-0337052P. 2001US-0348065P. 2001US-0336983P. 25-OCT-2002; 2002US-00282174. 2001US-0339525P 2002US-0368919P HOSPITAL CORP primer; ss; sequencing US2003224380-A1 Homo sapiens. 25-OCT-2001; 08-NOV-2001; 28-MAR-2002; 25-OCT-2001; 02-NOV-2001; 09-NOV-2001; 08-NOV-2001; 04-DEC-2001; 04-DEC-2003 (GEHO) GEN

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This invention relates to a novel method of determining a predisposition for or the occurrence of neurodegenerative disease comprising detecting in a target nucleic acid obtained from the subject the presence of an allelic variant of polymorphic regions of human genes selected from urokinase plasminogen activator (uPA), gamma-synuclein (SNCG), insulin lipase (LIPA) and tumour necrosis factor receptor SF6 (TNFRSF6). The method is useful in determining the presence or predisposition to a neurodegenerative disease, particularly Alzheimer's disease. The present sequence is that of a sequencing primer which was used for sequencing of a region of the human TNFRSF6 gene in the exemplification of the
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disease, particularly Alzheimer's disease, comprises determining the presence of a polymorphism in the uPA, SNCG, IDE, KNSL1, LIPA or TNFRSF6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetees; obesity; cardiovacular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2-methoxyethyl; 2'-MOB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human glucocorticoid receptor-specific antisense oligonucleotide #152.
                                                                                                                                                                                                                                                                                                                                                                                                               0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                             Example 3; SEQ ID NO 363; 205pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2214 ACAATGTGAGGGGTCC 2229
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 targets
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               the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       88;
                                                                                                                                                                                                                                                                                                                                                                                     Human glucocorticoid receptor-specific antisense oligonucleotide #136
                                                                                                                                                                                                                                                                                                                                                                                                                  antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obseity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
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present DNA sequence represents an antisense oligonucleotide that
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                                                                                                                                 1; Indels
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                                                                                            Score 14.4; DB 1; Length
Pred. No. 1.86+03;
0; Mismatches 1; Indels
                                                                Sequence 20 BP; 4 A; 0 C; 6 G; 10 T; 0 U; 0 Other;
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                                                                                                               93.8%;
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                                                                                                                                                                                                                                                                                   ADH63302 standard; DNA; 20
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                                                                                     Query Match
Best Local Similarity 93.89
Matches 15; Conservative
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Matches
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Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                    New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                            antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
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                                                                                        Human glucocorticoid receptor-specific antisense oligonucleotide #273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 4 A; 0 C; 6 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 273; 985pp; English
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                     ADH63439 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93.88;
                                                                 (first entry)
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                                                                                                                                                                                                                                                                                 (PHAA ) PHARMACIA CORP
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                                           ADH63439
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The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-NOE) wings and a phosphorothioate backbone.
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93.8%; Pred. No. 1.8e+03;
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Matches 15; Conservative
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                                                                                                                                                                                                                                                                                      (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-035034/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-035034/03.
WO2003099215-A2
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to nucleic acids encoding a mammalian glucocorticoid receptor. The antiesnes oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
             New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single nucleotide polymorphism in human gene.
                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.4; DB 1; Length 20; 13.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; single nucleotide polymorphism; SNP; ss; primer.
                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 5 A; 0 C; 6 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 8 A; 9 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; SEQ ID NO 5231; 2627pp; Japanese.
                                                                                         Claim 4; SEQ ID NO 782; 985pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                          3321 GAGATITATITIGG 3336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer of the invention #1922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAGATTTAGTTTTGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 93.8%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADK96202 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-SEP-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK96202;
                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 2069
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The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel polynucleotide useful for PCR amplification along with two fragment from another set of sequences, or for detecting single nucleotide polymorphism in human gene.
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0.4%; Score 14.4; DB 1; Length 20;
Best Local Similarity 93.8%; Pred. No. 1.8e+03;
Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interleukin; IL-4 receptor; IL-5 receptor; lung disease;
                                                                                                                                                                                                      human; single nucleotide polymorphism; SNP; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 9 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; SEQ ID NO 7057; 2627pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide associated to PDE4C #124.
                                                                                                                                                                                                                                                                                                                                                                                  (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                           Primer of the invention #3748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
                                                                                      ADK98028 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                          08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                                                                                                                                                     08-MAR-2002; 2002JP-00064373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ61058 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                19 TCTGTGCGTGTGTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         868 GAGGCTGACGAGGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16 checareaceaecee
WPI; 2004-093977/10.
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                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                             L6-SEP-2003
                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                   ADK98028;
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                                                                          ADK98028/
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Query Match 0.4%; Best Local Similarity 93.8%; Matches 15; Conservative

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0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.88+03; Ive 0; Mismatches 1; Indels

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uncleic acid molecule encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder; including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2,MOE wings and a deoxy gap. Used during the antisense inhibition of human Navi.3 expression, the oligonucleotides are designed to target different regions of the human Navi.3 RNA.
                                                                                                                          Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                     compound targeted to a nucleic acid molecule encoding
                                                                                                                                                                                                                                                                                            The present invention relates to an antisense compound targeted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric phosphorothioate oligonucleotide to target Nav1.3 #7343.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.4; DB 1; Length 20; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 5 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                            Claim 4; SEQ ID NO 948; 417pp; English.
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Best Local Similarity 93.8%;
Matches 15; Conservative
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ADK80009 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-203785/19.
                                                                       WPI; 2004-203785/19.
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                         Roberds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 2073
요
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-cend of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the cligonucleotide and optionally surfactant operatively linked to the cligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is conseful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded consent from airway inflammation, allergy(ies), asthma, impeded consent from airway inflammation, ung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                            Miller
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                                                                                                                                                                                            Nyce JW, Tang L, Sandrasagra A, Aguilar D,
Shahabuddin S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; SEQ ID NO 1914; 85pp; English
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                                             25-JUL-2003; 2003WO-US023509
                                                                                                29-JUL-2002; 2002US-0399076P
                                                                                                                                              (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CACGCAGGGCTGGCCC 16
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Matches 15; Conservative
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05-FEB-2004
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1; Indels

New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.

14-AUG-2003; 2003WO-US025465.

Synthetic.

ADK73614;

RESULT 2072

Query Match

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14-AUG-2002; 2002US-0403416P

(PHAA) PHARMACIA CORP

The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, pain from burns, migraine headache, cluster headache, mild-to-moderate beadache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with human Nav1.3 expression, the oligonucleotides inhibition of human Nav1.3 expression, the human Nav1.3 RNA. Query Match 0.4%; Score 14.4; DB 1; Length 20; Best Local Similarity 93.8%; Pred. No. 1.8e+03; Matches 15; Conservative 0; Mismatches 1; Indels Sequence 20 BP; 11 A; 1 C; 3 G; 5 T; 0 U; 0 Other; 4; SEQ ID NO 7343; 417pp; English 3261 ATATTTTATTTGCTTT 3276 14-AUG-2003; 2003WO-US025465 14-AUG-2002; 2002US-0403416P ß ADK73997 standard; DNA; 20 20 ATATTTTATTTGCCTT (first entry) (PHAA) PHARMACIA CORP WPI; 2004-203785/19. WO2004016754-A2. 20-MAY-2004 26-FEB-2004. Roberds SL; Synthetic ADK73997; RESULT 2074 Claim ADK73997

ADK73997

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Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss. Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1331.

New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.

The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3. e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate beadache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy, or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.

0.4%; Score 14.4; DB 1; Length 20;

Seguence 20 BP; 1 A; 6 C; 5 G; 8 T; 0 U; 0 Other;

Query Match

New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.

WPI; 2004-203785/19.

Claim 4; SEQ ID NO 1787; 417pp; English.

Claim 4; SEQ ID NO 1331; 417pp; English.

The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate

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headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navi.3 expression, the oligonucleotides are designed to target different regions of the human Navi.3 RNA.
                                                                                                                                                                                                                                                                                                                     Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                  Chimeric phosphorothioate oligonucleotide to target Navl.3 #1787.
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                                                                                                     0.4%; Score 14.4; DB 1; Length 20; 33.8%; Pred. No. 1.8e+03; ve 0; Mismatches 1; Indels
                                                                                 Sequence 20 BP; 1 A; 6 C; 4 G; 9 T; 0 U; 0 Other;
                                                                                                                                                      932
                                                                                                                                                                                                                                 ADK74453 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-AUG-2002; 2002US-0403416P.
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                                                                                                                  93.8%;
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                                                                                                                                                    917 TGGGCTTCTTCCTGTT
                                                                                                                                                                           TGAGCTTCTTCCTGTT
                                                                                                                  Local Similarity 93.8
nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                         Query Match
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Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.

14-AUG-2003; 2003WO-US025465 14-AUG-2002; 2002US-0403416P

WO2004016754-A2.

Synthetic.

26-FEB-2004.

(PHAA) PHARMACIA CORP

WPI; 2004-203785/19.

Roberds SL;

Chimeric phosphorothioate oligonucleotide to target Nav1.3 #5588.

(first entry)

20-MAY-2004

ADK78254;

ВЪ.

ADK78254 standard; DNA; 20

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                         Gaps
                                                                                                                                                                                                                                                                                                    Chimeric phosphorothioate oligonucleotide to target Navl.3 #3410.
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   93.8%; Pred. No. 1.8e+03; iive 0; Mismatches 1; Indels
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                                                           932
                                                                                                                                                                                        ADK76076 standard; DNA; 20 BP.
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                                                                                                4 TGAGCTTCTTCCTGTT 19
                                                                                                                                                                                                                                                                (first entry)
                                                         917 TGGGCTTCTTCCTGTT
                       15; Conservative
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                           ADK76076;
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New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or atexia.

The present invention relates to an antisense compound targeted to

Claim 4; SEQ ID NO 5588; 417pp; English

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Pred. No. 1.8e+03;
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93.8%; Pred. No. 1...
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Best Local Similarity 93.8
Matches 15; Conservative
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Gaps

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0.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 1.8e+03; ive 0; Mismatches 1; Indels

Query Match Best Local Similarity 93.8 Matches 15; Conservative

917 TGGGCTTCTTCCTGTT 932

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RESULT 2077 ADK78254

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nucleic acid molecule encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, lower back pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder; including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2,7MOE wings and a deoxy gap. Used during the antisense inhibition of human Navi.3 expression, the oligonucleotides are designed to target different regions of the human Navi.3 RNA.
                                                                                                                                  New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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14-AUG-2002; 2002US-0403416P.
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Beet Local Similarity 93.8%;
Conservative
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infantile epilepsy; ataxia;
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Matches 15; Conservative
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